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The invention describes an isolated secreted and transmembrane PRO polypeptide (I). PRO polypeptide such as PRO213, PRO700, PRO320 or PRO615 or use useful in biotechnological and medical research, as well as in various industrial applications. PRO polypeptide such as PRO300, PRO302, PRO932, PRO9120, PRO9120,
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Human; ss; PCR; secreted protein; transmembrane protein; PRO; cytostatic; ophthalmological; antiarchritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth, retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer. 2.9%; Score 12.4; DB 1; Length 18; 92.9%; Pred. No. 4.3e+02; rive 0; Mismatches 1; Indels ADC68610 standard; DNA; 18 BP. Human PRO 274 PCR primer #4 GAACTCGGTGGCGG 228 18-DEC-2003 (first entry) Best Local Similarity 92.9 Matches 13; Conservative US2003064407-A1. Homo sapiens. 03-APR-2003.

ADC68610;

RESULT 698 ADC686

Query Match

215 8 97US-0064249P 97US-0065411P 97US-0065311P 98US-0077450P 98US-0077632P 98US-0077641P 98US-0077641P

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03-NOV-1997; 13-NOV-1997; 21-NOV-1997;

17-OCT-1997;

24-OCT-2001; 2001US-00999834

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98US-0080165P.
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98US-0081817P.
98US-0081819P.
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15-MAY-1998; 98US-0086704P.
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26-JUN-1998; 98US-008720P.
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28-MAY-1999; 99US-0113621P.
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22-DEC-1999; 99US-0113621P.
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23-DEC-1999; 99US-0113621P.
23-DEC-1999; 99US-0113621P.
23-DEC-1999; 99US-0113621P.
23-JUN-1999; 99US-0113621P.
24-MAR-1999; 99US-0113621P.
25-AUG-1999; 99US-0113621P.
26-JUN-2000; 2000WO-US001365.
11-FEB-2000; 2000WO-US001361.
11-FEB-2000; 2000WO-US001361.
11-FEB-2000; 2000WO-US001361.
11-FEB-2000; 2000WO-US001361.
11-MAR-2000; 2000WO-US001361.
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Human, 88; PCR; secreted protein, transmembrane protein, PRO; oytostatic, ophthalmological, antiarthritic, osteopathic; antirheumatic; vulnerary; auditory, tumour growth, retinal disorder; sports-related joint problem, articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.
                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                   Desnoyers L, Eaton DL;
                                                                                                                                                                                                                                                                        Length 18;
                                                                                                                                                                                                                                                                       Query Match
2.9%; Score 12.4; DB 1;
Best Local Similarity 92.9%; Pred. No. 4.3e+02;
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01-DEC-2000, 2000Wo-US032678.
20-DEC-2000; 2000WS-00747259.
20-DEC-2000; 2000WS-00747259.
28-PEB-2001; 2001WS-00806520.
22-MAR-2001; 2001WS-00816744.
22-MAR-2001; 2001WS-00816744.
22-MAR-2001; 2001WS-00816749.
22-MAR-2001; 2001WS-00854280.
10-MAY-2001; 2001WS-00854280.
25-MAY-2001; 2001WS-00854280.
25-MAY-2001; 2001WS-00872035.
01-UNN-2001; 2001WS-00872035.
01-UNN-2001; 2001WS-00882636.
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39-UUL-2001; 2001WS-0US019692.
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970S-0064249P.
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ADC62670 standard; DNA; 18
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                                                                                                                                                                                                                              (GETH ) GENENTECH INC.
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03-NOV-1997;
13-NOV-1997;
10-MAR-1998;
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31-MAR-1998;
32-MAR-1998;
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07-MAY-1998
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26-JUN-1998; 98US-0090863P.
26-JUN-1998; 98US-0091010P.
30-JUL-1998; 98US-0091010P.
30-JUL-1998; 98US-009163EP.
30-JUL-1998; 98US-010038P.
11-SEP-1998; 98US-010038P.
20-NOV-1998; 98US-010034FP.
22-NOV-1998; 98US-010034FP.
23-DEC-1998; 98US-010324FP.
23-DEC-1999; 99US-013285P.
24-ARR-1999; 99US-013023EP.
25-ARR-1999; 99US-013023EP.
26-ARR-1999; 99US-013023EP.
26-ARR-2009; 90US-013023EP.
26-ARR-2000; 2000WO-US002312.43.
30-DEC-1999; 99WO-US003212.43.
30-DEC-1999; 99WO-US CGXSX

(GETH) GENENTECH INC

Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL;
Stewart TA, Tumas D, Williams PM, Wood WI;

WPI; 2003-695924/66.

New isolated secreted and transmembrane PRO polypeptides, useful in the preparation of a medicament for treating a condition responsive to the polypeptide, and as therapeutic agents e.g. vaccines.

Example 4; SEQ ID NO 14; 467pp; English.

The invention relates to an isolated PRO polypeptide (secreted or transmembrane protein) having at least 80% amino acid sequence identity

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to an amino acid sequence chosen from 94 fully defined sequences as given in the specification (including PRO lacking its associated signal peptide, a PRO extracellular domain with or without its associated signal peptide. Also included are neuclaic acids encoding the PRO proteins mentioned above, a vector comprising a PRO nucleic acid), a host cell comprising the vector and producing PRO, a chimaeric molecule comprising PRO fused to a heterologous amino acid sequence, and an anti-PRO antibody. PRO337 polypeptide is useful for detecting a PRO4993 antibody. PRO4993 polypeptide in a sample suspected of containing PRO4993 polypeptide in a sample suspected of containing PRO4993 polypeptide, and PRO159 polypeptide is useful for detecting PRO155 polypeptide, and PRO159 polypeptide is useful for detecting PRO155 PRO700 or PRO739. PRO4993 polypeptide is useful for linking a ploantie is the toxin, radiolabel, or an antibody. The bloactive molecule is the cell. PRO337 polypeptide is useful for linking a causes death of the cell. PRO337 polypeptide is useful for linking a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; ss; PCR; secreted protein; transmembrane protein; PRO; cytostatic; ophthalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth; retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.
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                                                                                                                                                                                                                                    Query Match 2.9%; Score 12.4; DB 1; Length 18; Best Local Similarity 92.9%; Pred. No. 4.3e+02; Matches 13; Conservative 0; Mismatches 1; Indels
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03-NOV-1997;
21-NOV-1997;
21-NOV-1997;
10-MAR-1998;
11-MAR-1998;
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11-MAR-1998;
12-MAR-1998;
20-MAR-1998;
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20-MAR-1998;
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26 - MAR - 1998;
27 - MAR - 1998;
30 - MAR - 1998;
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PR 31-MR-1996 PB 98US-0080105P.
PR 01-ARF-1998 PB 08US-0080134P.
PR 01-ARF-1998 PB 08US-0080134P.
PR 01-ARF-1998 PB 08US-0080134P.
PR 01-ARF-1998 PB 08US-0080134P.
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Human, 88; PCR; secreted protein; transmembrane protein; PRO; cytostatic; ophthalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth; retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.
peptide). Also included are nucleic acids encoding the PRO proteins mentioned above, a vector comprising a PRO nucleic acid), a host cell comprising the vector and producing PRO, a chimaeric molecule comprising PRO fused to a heterologous amino acid sequence, and an anti-PRO antibody. PRO337 polypeptide is useful for detecting a PRO4993 polypeptide is useful for detecting a PRO4993 polypeptide is useful for detecting PRO337 polypeptide. Similarly, PRO4993 polypeptide is useful for detecting PRO359 polypeptide, and PRO1559 polypeptide is useful for detecting PRO1559 polypeptide is useful for detecting
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2.9%; Score 12.4; DB 1; Length 18;
Best Local Similarity 92.9%; Pred. No. 4.3e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
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ADC41055 standard; DNA; 18
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03-NOV-1997;
21-NOV-1997;
11-MAR-1998;
11-MAR-1998;
11-MAR-1998;
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20-MAR-1998;
20-MAR-1998;
21-MAR-1998;
27-MAR-1998;
27-MAR-1998;
27-MAR-1998;
31-MAR-1998;
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01-APR-1998;
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ADC41055/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
Goddowski PJ, Grimaldi JC, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton
Stewart TA, Tumas D, Williams PM, Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel secreted and transmembrane polypeptides, designated PRO polypeptides, and polynucleotides encoding them useful for treating kidney diseases, bone, cartilage and retinal disorders.
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       22-DEC-1998, 98US-0113296P.

23-DEC-1998, 98US-0113621P.

06-JAN-1999, 99WO-USO00106.

08-MAR-1999, 99WO-USO0028.

12-MAR-1999, 99WO-USO05190.

12-MAR-1999, 99WS-0126773P.

25-MAR-1999, 99US-0130222P.

26-APR-1999, 99US-0131445P.

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14-MAY-1999, 99US-0131445P.

14-MAY-1999, 99US-0131445P.

14-MAY-1999, 99US-013722P.

28-JUL-1999, 99US-0139557P.

29-CCT-1999, 99WS-01802252.

29-CCT-1999, 99WS-018028551.

20-DEC-1999, 99WS-018028551.

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24-JUL-1999, 99WS-018028551.

25-CCT-1999, 99WS-018028551.

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26-CCT-1999, 99WS-0180281.

27-CCT-1999, 99WS-0180281.

28-CCT-1999, 99WS-018031.

29-CCT-1999, 99WS-018031.

24-FEB-2000, 2000WS-01801.

24-FEB-2000, 2000WS-01801.

25-MAX-2000, 2000WS-01801.

26-CUN-2000, 2000WS-01801.

27-MAX-2000, 2000WS-01801.

28-CUN-2000, 2000WS-01808.

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98US-0081953P.
98US-0081952P.
98US-0081953P.
98US-0081953P.
98US-0081953P.
98US-0081953P.
98US-0081953P.
98US-0081953P.
98US-0081953P.
98US-0081953P.
98US-0081953P.
98US-008193P.
98US-008198P.
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99WO-US005028.
99WO-US005190.
01-APR-1998;

08-APR-1998;

08-APR-1998;

09-APR-1998;

15-APR-1998;

15-APR-1998;

15-APR-1998;

15-APR-1998;

15-APR-1998;

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07-MAY-1998;

07-MAY-1998;

07-MAY-1998;

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07-MAY-1998;
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13-MAX-1998;

13-MAX-1998;

15-MAX-1998;

15-MAY-1998;

15-MAY-1998;

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22-MAY-1998;

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23-MAY-1998;

24-MAY-1998;

28-MAY-1998;

28-MAY-1998;

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28-MAY-1998;

28-MAY-1998;
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11-SEP-1998;
07-OCT-1998;
20-NOV-1998;
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12-MAR-1999; 99US-0123957P.
29-MAR-1999; 99US-01205673P.
21-APR-1999; 99US-0131022P.
26-APR-1999; 99US-0131022P.
26-APR-1999; 99US-0131022P.
26-AUX-1999; 99US-0131445P.
26-UUN-1999; 99WO-US010733.
23-UUN-1999; 99WO-US010733.
23-UUN-1999; 99WO-US010737P.
23-UUN-1999; 99WO-US010737P.
26-UUL-1999; 99WO-US01257P.
26-UUL-1999; 99WO-US01257P.
26-UUL-1999; 99WO-US012813.
26-UUL-1999; 99WO-US012813.
26-UUL-1999; 99WO-US012813.
28-UUL-1999; 99WO-US018137P.
28-UUL-1999; 99WO-US01813.
29-DEC-1999; 99WO-US01813.
21-DEC-1999; 99WO-US01813.
23-UNN-2000; 2000WO-US01095.
21-WAR-2000; 2000WO-US01095.
21-WAR-2000; 2000WO-US01319.
21-WAR-2000; 2000WO-US01319.
21-WAR-2000; 2000WO-US01319.
21-WAR-2000; 2000WO-US01813.
21-WAR-2000; 2000WO-US01319.
21-WAR-2000; 2000WO-US01319.
21-WAR-2000; 2000WO-US01319.
22-WAY-2000; 2000WO-US01319.
23-WAY-2000; 2000WO-US01319.
24-WUN-2000; 2000WO-US013189.
25-WAY-2000; 2000WO-US013189.
25-WAY-2001; 2011WO-US01318.
25-WAY-2001; 2011WO-US01318.
25-UNN-2001; 2011WO-US01318.
25-UNN-2001; 2011WO-US01318.
25-UNN-2001; 2011WO-US01318.
25-UNN-2001; 2011WO-US01318.
25-UNN-2001; 2011WO-US01318.

(GETH) GENENTECH INC

Ashkenazi A, Baker KP, Botstein D, Desnoyers L, Eaton DL;
Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL;
Stewart TA, Tumas D, Williams PM, Wood WI;

WPI; 2003-743806/70.

Novel isolated secreted and transmembrane PRO polypeptides, useful in the preparation of a medicament for treating a condition responsive to the polypeptide, and as therapeutic agents e.g. vaccines.

Example 4; SEQ ID NO 14; 466pp; English.

The invention relates to an isolated PRO polypeptide (secreted or transmembrane protein) having at least 80% amino acid sequence identity to an amino acid sequence chosen from 94 fully defined sequences as given in the specification (including PRO lacking its associated signal peptide, a PRO extracellular domain with or without its associated signal peptide). Also included are nucleic acids encoding the PRO proteins entioned above, a vector comprising a PRO nucleic acid), a host cell comprising the vector and producing PRO, a chimaeric molecule comprising PRO fused to a heterologous amino acid sequence, and an anti-PRO antibody. PRO337 polypeptide is useful for detecting a PRO4993

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98US-00812292-
98US-00818179-
98US-008181179-
98US-0081818P-
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98US-0109304P.
98WO-US024855.
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98US-0113621P.
99WO-US000106.
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99US-0131022P.
99US-0131445P.
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22-MAY-1998;
22-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                        28-MAY-1998
ö
                                                                                                                                                                     vulnerary; virucide; neuroprotective; cytostatic; gene therapy; tumour cell proliferation inhibitor; secereted and transmembrane protein; PRO; viral infection; wound healing; tissue growth; muscle generation; muscle regeneration; amyotrophic lateral sclerosis; neuropathy; AlDS-associated neuropathy; diabetic peripheral neuropathy; chromosome identification; antagonist; tissue typing; immunohistochemical staining; primer; ss.
                                     Gaps
polypeptide in a sample suspected of containing PRO4993 polypeptide.
Similarly, PRO4993 polypeptide is useful for detecting PRO337
                                      ö
                     Query Match 2.9%; Score 12.4; DB 1; Length 18; Best Local Similarity 92.9%; Pred. No. 4.3e+02; Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                 97US-0062250P.
97US-0064249P.
97US-0065311P.
98US-0077450P.
98US-0077641P.
98US-0077641P.
98US-0077641P.
98US-0077641P.
                                                                                                                                                                                                                                                                                 25-OCT-2001; 2001US-00016177
                                                                                                                                                        Human PRO 274 PCR primer #4.
                                                      215 GAACTCGGTGGCGG 228
                                                                                                   110/c
ADC67110 standard; DNA; 18
                                                                                                                                         18-DEC-2003 (first entry)
                                                                GAACTCCGTGGCGG 5
                                                                                                                                                                                                                                                   US2003073131-A1.
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27-MAR-1998;
27-MAR-1998;
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08-APR-1998;
09-APR-1998;
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30-MAR-1998;
31-MAR-1998;
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31-MAR-1998
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                                                                                                                         ADC67110;
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                                                                                            RESULT 702
                                                                                                    ADC67110/
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Example 4; SEQ ID NO 14; 464pp; English.
14-WAY-1999; 99US-0134287P.
14-WAY-1999; 99WO-USO11733.
02-JUN-1999; 99WO-USO11735.
16-JUN-1999; 99WO-USO1139557P.
23-JUN-1999; 99WS-013139557P.
23-JUN-1999; 99US-0142680P.
26-JUL-1999; 99US-0142680P.
26-JUL-1999; 99US-0142680P.
28-JUL-1999; 99US-0142622P.
29-OCT-1999; 99US-0142622P.
29-OCT-1999; 99WO-USO18313.
02-DEC-1999; 99WO-USO18253.
16-DEC-1999; 99WO-USO1825.
16-DEC-1999; 99WO-USO1825.
16-DEC-1999; 99WO-USO1824.
16-DEC-1999; 99WO-USO1824.
11-FEB-2000; 2000WO-USO0057.
11-FEB-2000; 2000WO-USO0057.
11-FEB-2000; 2000WO-USO01841.
11-MAR-2000; 2000WO-USO01841.
11-MAR-2000; 2000WO-USO01841.
11-MAR-2000; 2000WO-USO18705.
21-MAR-2000; 2000WO-USO18705.
21-MAR-2000; 2000WO-USO18705.
21-MAR-2000; 2000WO-USO18705.
22-MAR-2000; 2000WO-USO18705.
23-MAR-2000; 2000WO-USO18705.
24-MUG-2000; 2000WO-USO18705.
25-MAR-2001; 2001WO-USO19692.
25-MAR-2001; 2001WO-USO19692.
25-MAR-2001; 2001WO-USO11965.
25-MAR-2001; 2001WO-USO11968.
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Novel isolated secreted and transmembrane PRO polypeptides, useful in t preparation of a medicament for treating a condition responsive to the polypeptide, and as therapeutic agents e.g. vaccines.
Ashkenazi AJ, Baker KD, Botstein D, Desnoyers L, Eaton DL;
Ferrara N, Filvaroff B, Fong S, Gao W, Gerber H, Gerritsen ME;
Goddard A, Goddwski PJ, Grimaldi JC, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton Stewart TA, Tumas D, Williams PM, Wood WI;
```

The invention describes an isolated secereted and transmembrane PRO polypeptide (1). PRO polypeptide such as PRO213, PRO700, PRO320 or PRO615 is useful in biotechnological and medical research, as well as in various industrial applications. PRO polypeptide such as PRO300, PRO566, PRO703, PRO560 or PRO646, PRO351, PRO52, PRO381, PRO615, PRO616, PRO703, PRO660 or PRO646 is useful for therapeutic purposes. PRO363 is useful therapeutically in vivo for lessening the effects of viral infection. PRO200 is useful for the treatment of wound healing, tissue growth and muscle generation and regeneration. PRO337 is useful for treating amyotrophic lateral sclerosis, neuropathy, AlDS-associated neuropathy or

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                                                     Gaps
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Ouery Match
2.9%; Score 12.4; DB 1; Length 18;
Best Local Similarity 92.9%; Pred. No. 4.3e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
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Human; ss; PCR; secreted protein; transmembrane protein; PRO; cytostatic; ophthalmological; antiarchritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth; retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.
                                                                                                                                                                                                                                 97US - 0062250P.
97US - 006424PP.
97US - 00634EP.
98US - 0077450P.
98US - 0077641P.
98US - 007804P.
98US - 007804P.
98US - 007804P.
98US - 007804P.
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9805-0079788F

9805-0079920F

9805-0079923F

9805-0080105F

9805-0080107F

9805-0080337F

9805-0080333F

9805-0080333F

9805-0080333F

9805-0080333F
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98US-0081203P.
98US-0081229P.
98US-0081817P.
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98US-0082568P.
98US-0082569P.
                                                                                                                                                                                                                  15-OCT-2001; 2001US-00978193
                                                                                                     Human PRO 274 PCR primer #4
GAACTCGGTGGCGG 228
                                                      ADC62046 standard; DNA; 18
                                                                                     18-DEC-2003 (first entry)
             GAACTCCGTGGCGG 5
                                                                                                                                                                                  US2003073624-A1.
                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                   17-APR-2003
                                                                     ADC62046;
215
              8
                                        RESULT 703
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98US-0082700P.
98US-0082704P.
98US-0082704P.
98US-008322P.
98US-0083322P.
98US-0083495P.
98US-0083322P.
98US-0083495P.
98US-0083495P.
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98US-0083495P.
98US-0083495P.
98US-0083495P.
98US-0083495P.
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98US-0084637P.
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98US-0084643P.
98US-0084643P.
98US-008559P.
98US-008569P.
98US-001038P.
99US-0013637P.
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07-MAY-1998,
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07-MAY-1998,
13-MAY-1998,
13-MAY-1998,
13-MAY-1998,
15-MAY-1998,
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15-MAY-1998,
15-MAY-1998,
15-MAY-1998,
15-MAY-1998,
16-MAY-1998,
18-MAY-1998,
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14-MAY-1999; 9908-00311832.
14-MAY-1999; 9908-01311832.
14-MAY-1999; 9908-01311832.
14-MAY-1999; 9908-01311832.
15-UJN-1999; 9908-0145680P.
28-UJL-1999; 9908-0145680P.
28-MJC-1999; 9908-0145680P.
28-MJC-1999; 9908-01380138.
28-MJC-1999; 9908-01380138.
29-OCT-1999; 9908-0162506P.
20-DEC-1999; 9908-0162506P.
20-DEC-2000; 2000WO-US00143941.
20-DEC-2000; 2000WO-US

Query Match
2.9%; Score 12.4; DB 1;
Best Local Similarity 92.9%; Pred. No. 4.3e+02;
Matches 13; Conservative 0; Mismatches 1; GAACTCGGTGGCGG 228 215 18

(GETH) GENENTECH INC.

Length 18;

RESULT 704 ADC13477/c

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US2003104998-A1.
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11 - MAR - 1998 |
12 - MAR + 1998 |
13 - MAR - 1998 |
17 - MAR - 1998 |
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20 - MAR - 1998 |
21 - MAR - 1998 |
22 - MAR - 1998 |
27 - MAR - 1998 |
28 - MAR - 1998 |
29 - MAR - 1998 |
20 - MAR - 1998 |
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30 - MAR - 1998 |
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08-APR-1998;
08-APR-1998;
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15-APR-1998;
15-APR-1998;
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31-MAR-1998;
31-MAR-1998;
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01-APR-1998;
01-APR-1998;
01-APR-1998;
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15-APR-1998;
15-APR-1998;
                                                                                                                                                                                               Homo sapiens.
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09-APR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a novel method for determining whether a treatment is effective in changing a status of a certain set of target cells in an individual. The method comprises obtaining a sample from an individual after initiation of the treatment; and determining whether the sample comprises an expression product of at least one marker gene. The marker gene and a proteinaceous molecule (which can bind to the protein derived from the marker gene of the invention) are useful for determining whether a treatment is effective in counteracting a tumour in an individual, especially Kaposi's Sarcoma. Peripheral blood mononuclear cell (PBMC) expressed keratin 14, TIE 1, Salioadheain, or Siglec 1 sequences or a fully defined sequence given in the specification. The their analogues are useful as indicators for angiogenesis and for their analogues comprising a marker gene of the invention is useful as a drug target. The compound is useful for preparing a medicament. This polynuclectide sequence represents a PCR primer of a Kaposi's Sarcoma tag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Determining whether a treatment is effective in changing a status of a certain set of target cells in an individual comprises determining whether the sample comprises an expression product of at least one marker
                                                                                                                                                                                                                marker gene; tumour; Kaposi's Sarcoma; peripheral blood mononuclear cell; PBMC; expressed keratin 14; TIE 1; Salioadhesin; Siglec 1; angiogenesis; drug target; tag; SAGE library; KS3; KS4; PCR; primer; ss.
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Pred. No. 4.3e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 2 A; 8 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                              Kaposi's sarcoma tag PCR primer, SEQ ID No 144.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 144; 94pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-SEP-2001; 2001EP-00203703.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (PRIM-) PRIMAGEN HOLDING BV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 288 AAGCTGGTGAAGGA 301
ADC13477 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18 AAGCTGCTGAAGGA S
                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-589342/56.
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                                                                                                                                                                                                                                                                                                                                                                                     EP1298221-A1
                                                                                                                                                                                                                                                                                                                                   Unidentified
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                                                                                                            18-DEC-2003
                                                    ADC13477;
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Matches
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ADC41679 standard; DNA; 18 BP

ADC41679,

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Human PRO 274 PCR primer #4

18-DEC-2003

8X2X5X8

ADC41679;

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Human, 88; PCR; secreted protein, transmembrane protein; PRO; cytostatic; ophthalmological; antiarchritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth; retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer
                                                                                                                        97US-0064249P.
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97US-0065864P.
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98US-0079264P.
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98US-0080333P.
98US-0081265P.
98US-0081265P.
98US-008137P.
                                                                                                          16-OCT-2001; 2001US-00978643
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27-APR-1998;
28-APR-1998;
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29-OCT-1999; 99US-0162506P.
30-NOV-1999; 99W0-US028313.
30-DEC-1999; 99W0-US028551.
02-DEC-1999; 99W0-US028551.
16-DEC-1999; 99W0-US0310245.
30-DEC-1999; 99W0-US0310245.
30-DEC-1999; 99W0-US0310245.
30-DEC-1999; 99W0-US0310243.
30-DEC-1999; 99W0-US0310243.
30-DEC-1999; 99W0-US0310243.
30-DEC-1999; 99W0-US0310243.
30-DEC-1999; 99W0-US0310243.
31-FEB-2000; 2000W0-US000219.
05-JAN-2000; 2000W0-US0003765.
31-FEB-2000; 2000W0-US000376.
31-FEB-2000; 2000W0-US000376.
31-FEB-2000; 2000W0-US000376.
31-FEB-2000; 2000W0-US000376.
31-FEB-2000; 2000W0-US000376.
32-MAX-2000; 2000W0-US003238.
33-MAX-2000; 2000W0-US013238.
34-FEB-2000; 2000W0-US013238.
35-MAX-2000; 2000W0-US013238.
37-NOV-2000; 2000W0-US013238.
37-NOV-2000; 2000W0-US013264.
38-MAX-2001; 2001W0-US0132678.
39-MAX-2001; 2001W0-US013682.
32-MAX-2001; 2001W0-US013683.
32-MAX-2001; 2001W0-US013683.
32-MAX-2001; 2001W0-US013683.
33-UUN-2001; 2001WS-US013683.
30-UUN-2001; 2001WS-US01368.
30-UUN-2001; 2001WS-US01368.
30-UUN-2001; 2001WS-US01368.
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     98US-0083554P.
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98US-0083558P.
98US-0083358P.
98US-0084341P.
98US-0084411P.
98US-0084538P.
98US-0084637P.
98US-0084637P.
98US-0084637P.
98US-0084637P.
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99US-01314287P.
99US-01314287P.
       29-APR-1998;
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30-APR-1998;
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30-APR-1998;
66-MAY-1998;
66-MAY-1998;
67-MAY-1998;
67-MAY-1999;
67-MAY-1998;
67-MA
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Query Match
Best Local Similarity 92.9%; Pred. No. 4.30+02;
Matches 13; Conservative 0; Mismatches 1; Indels 0; Gaps 0

215 GAACTCGGTGGCGG 228

CY 215 GAACTCCGTGGCGG 5

BESULT 706

AD 49048 Standard; DNA; 18 BP.

XX
AC ADE49048 Standard; DNA; 18 BP.

XX
AC ADE49048;

XX
AC ADE49048;

XX
AC ADE49048;

XX
AC ADE49048;

XX
AC ADE40048;

AC ADE4
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98US-0086313P.
98US-0085313P.
98US-008563P.
99US-008563P.
99US-008563P.
99US-008643P.
99US-008663P.
07-MAY-1998,
07-MAY-1998,
07-MAY-1998,
07-MAY-1998,
07-MAY-1998,
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13-MAY-1998,
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15-MAY-1998,
16-MAY-1998,
22-MAY-1998,
22-MAY-1998,
22-MAY-1998,
22-MAY-1998,
22-MAY-1998,
22-MAY-1998,
26-UUN-1998,
01-UUL-1998,
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05-JAN-2000;
    97105-0064249P.
97105-0064449P.
97105-00665644P.
97105-00665644P.
97105-00665644P.
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98105-00776649P.
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98105-0077664P.
98105-00779786P.
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98105-0081203P.
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98105-0081203P.
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98105-0081302P.
98105-0081302P.
98105-0081302P.
98105-0081316P.
98105-0081816P.
                                                                                                                                2002US-00978187
                                              US2003096744-A1
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03-NOV-1997;

21-NOV-1997;

21-NOV-1997;

11-MAR-1998;

11-MAR-1998;

11-MAR-1998;

12-MAR-1998;

12-MAR-1998;

20-MAR-1998;

20-MAR-1998;

20-MAR-1998;

20-MAR-1998;

27-MAR-1998;

27-MAR-1998;

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27-MAR-1998;

30-MAR-1998;

31-MAR-1998;

31-MAR-1998;
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09-APR-1998;
15-APR-1998;
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15-APR-1998;
15-APR-1998;
15-APR-1998;
21-APR-1998;
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39-APR-1998
39-APR-1998
39-APR-1998
39-APR-1998
39-APR-1998
39-APR-1998
30-APR-1998
        Homo sapiens
                                                                                                                            28-JAN-2002;
                                                                                      22-MAY-2003
      X_{X} \times X_{X
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Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
Goddard A, Godowski PJ, Grimaldi UC, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton I
Stewart TA, Tumas D, Williams PM, Wood WI;

15-MAY-1998; 98US-0085689F.
08-MAR-1999; 99WG-US005028.
28-ARR-1999; 99US-0131445F.
25-AUG-1999; 99US-00380138.
18-FEB-2000; 2000WG-US004341.
30-UUL-2001; 2001US-00918585.

(GETH) GENENTECH INC.

New genes, and its encoded secreted and transmembrane polypeptides, useful for treating e.g. lung or breast tumors, osteoarthritis, rheumatoid arthritis, obesity, diabetes, hyperinsulinemia, hypoinsulinemia or wounds.

WPI; 2003-875641/81.

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Human, ss, PCR, secreted protein; transmembrane protein, PRO; cytostatic, ophthalmological, antiarthritic; osteopathic; antirheumatic; vulnerary; auditory, tumour growth, retinal disorder; sports-related joint problem, articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 2.9%; Score 12.4; DB 1; Length 18; Best Local Similarity 92.9%; Pred. No. 4.3e+02; Matches 13; Conservative 0; Mismatches 1; Indels
06-JAN-2000; 2000WO-US000277.
06-JAN-2000; 2000WO-US000376.
11-FEB-2000; 2000WO-US000376.
12-FEB-2000; 2000WO-US00565.
13-FEB-2000; 2000WO-US005641.
10-MAR-2000; 2000WO-US0056319.
11-MAR-2000; 2000WO-US0056319.
12-MAR-2000; 2000WO-US0013705.
13-MAR-2000; 2000WO-US013705.
13-MAR-2000; 2000WO-US013705.
13-MAR-2000; 2000WO-US013705.
13-MAR-2000; 2000WO-US013705.
14-MAR-2000; 2000WO-US013705.
15-MAR-2000; 2000WO-US01564.
16-MAR-2000; 2000WO-US01564.
16-MAR-2000; 2000WO-US01566.
16-MAR-2000; 2000WO-US0156.
16-MAR-2001; 2000WO-US032678.
16-MAR-2001; 2000WO-US036520.
10-MAR-2001; 2000WO-US016920.
10-MAR-2001; 2000WO-US016920.
10-MAR-2001; 2000WO-US016920.
10-MAR-2001; 2000WO-US016920.
10-MAR-2001; 2000WO-US016920.
10-MAR-2001; 2000WO-US017680.
10-MAR-2001; 2000WO-US017680.
11-UWN-2001; 2000WO-US017680.
11-UWN-2001; 2000WO-US01765.
11-UWN-2001; 2000WO-US01765.
11-UWN-2001; 2000WO-US01765.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   215 GAACTCGGTGGCGG 228
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADE35102 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-JAN-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GETH ) GENENTECH INC.
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ADE35102/c
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The invention relates to an isolated PRO polypeptide (secreted or transmembrane protein) having at least 80% amino acid sequence identity to an amino acid sequence chosen from 94 fully defined sequences as given in the specification (including PRO lacking its associated signal peptide, a PRO extracellular domain with or without its associated signal comprising a PRO acids encoding the PRO proteins mentioned above, a vector comprising a PRO nucleic acid), a host cell comprising the vector and producing PRO, a chimaeric molecule comprising prolypeptide is useful for detecting a PRO4993 polypeptide is useful for inking a bioactive molecule to a cell expressing PRO4993 polypeptide is useful for inking a bioactive molecule to a cell expressing PRO4993 polypeptide is useful for inking a bioactive molecule to a cell expressing PRO4993 polypeptide is useful for inking a bioactive molecule to a cell expressing PRO4993 polypeptide is useful for inking a bioactive molecule to a cell expressing PRO4993 polypeptide is useful for modulating the polypeptide or molecule to a cell expressing PRO4993 polypeptide is useful for modulating at least one biological activity of the cell expressing PRO4993 polypeptide is useful for modulating the biological activity of the cell expressing PRO4993 polypeptide or an anti-PRO493 polypeptide or an anti-PRO493 polypeptide or modulating the biological activity of the cell expressing PRO4993 polypeptide or an anti-PRO493 polypeptide or an anti-PRO493 polypeptide or modulating the biological activity of the cell expressing PRO4993 polypeptide or an anti-PRO493 polypeptide or an anti-PRO4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 4; SEQ ID NO 14; 462pp; English.
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ADE16216 standard; DNA; 18 BP.
           Human PRO 274 PCR primer #4
        29-JAN-2004 (first entry)
      ADE16216;
RESULT 708
 ADE16216/
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Human; ss; PCR; secreted protein; transmembrane protein; PRO; cytostatic; obthchalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth; retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.

Homo sapiens.

US2003203435-A1.

30-OCT-2003.

18-OCT-2001; 2001US-00145092,

30-APR-1998; 98US-0083742P. 08-MAR-1999; 99WG-US005028. 23-JUN-1999; 99US-0141037P. 25-AUG-1999; 99US-00380138. 18-FBB-2000; 2000WG-US004341. 30-JUL-2001; 2001US-00918585.

(GETH) GENENTECH INC.

Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL; Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME; Goddard A, Godwski PJ, Grimaldi JC, Gurney AL, Hillan KJ; Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL; Stewart TA, Tumas D, Williams PM, Wood WI;

WPI; 2003-875642/81.

New genes, and its encoded secreted and transmembrane polypeptides, useful for treating e.g. lung or breast tumors, osteoarthritis, rheumatoid arthritis, obesity, diabetes, hyperinsulinemia, hypoinsulinemia or wounds.

Example 4; SEQ ID NO 14; 452pp; English.

The invention relates to an isolated PRO polypeptide (secreted or transmembrane protein) having at least 80% amino acid sequence identity con an amino acid sequence chosen from 94 fully defined sequences as given in the specification (including PRO lacking its associated signal peptide, a PRO extracellular domain with or without its associated signal comprising the vector comprising a PRO nucleic acid), a host cell comprising the vector and producing PRO, a chimaeric molecule comprising comprising the vector and producing PRO, a chimaeric molecule comprising comprising the vector and producing PRO, a chimaeric molecule comprising anti-body. PRO337 polypeptide is useful for detecting a PRO4993 polypeptide is useful for detecting a PRO4993 polypeptide is useful for detecting pRO375. PRO700 or PRO795 polypeptide is useful for detecting pRO375. PRO700 or PRO795 polypeptide is useful for detecting a pRO1559 polypeptide is useful for detecting a pRO355, PRO700 or PRO793 polypeptide is useful for inking a bloactive molecule to a cell expressing PRO337 polypeptide. The bloactive molecule to a cell expressing PRO493 polypeptide; PRO725, PRO700 or PRO739 polypeptide; as useful for linking a bloactive molecule to a cell expressing PRO493 polypeptide; professing PRO755, PRO700 or PRO739 polypeptide; and PRO1559 polypeptide; professing PRO755, PRO700 or PRO739 polypeptide; and PRO1559 polypeptide is useful for linking a bloactive molecule to a cell expressing PRO4933 polypeptide; professing PRO755, PRO700 or PRO739 polypeptide are useful for linking a bloactive molecule to a cell expressing PRO4933 polypeptide; professing PRO755, PRO700 or PRO739 polypeptide. DRO4993 polypeptide or anti-PRO337 polypeptide or anti-PRO337 polypeptide or anti-PRO3337 p

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the cell expressing PRO337 polypeptide, where the cell is killed. PRO337 polypeptide is useful for modulating the bollogical activity of the cell expressing PRO4939 polypeptide is useful for modulating the PRO700 or PRO739 polypeptide or an anti-PRO1559 polypeptide is useful for modulating the biological activity of the cell expressing PRO155 polypeptide, and PRO1559 polypeptide, and PRO1559 polypeptide or anti-PRO7025, anti-PRO700 or anti-PRO739 polypeptide, serving for anti-PRO700 or anti-PRO739 polypeptide is useful for modulating the biological activity of the cell expressing PRO725, proform or PRO739 polypeptide. The polypeptides are useful for inhibiting tumour growth, retinal disorders, sports-calated joint problems, articular cartilage defects, osteoarthritis or rheumatoid arthritis, wound healing and hearing loss in amammals. The present sequence is a PCR primer used to isolate nucleic acid encoding a PRO protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New PRO genes and encoded secreted and transmembrane polypeptides, useful for treating e.g. lung or breast tumors, osteoarthritis, rheumatoid arthritis, obesity, diabetes, hyperinsulinemia, hypoinsulinemia or wounds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human, 88; PCR; secreted protein, transmembrane protein, PRO, cytostatic, ophthalmological, antiarthritic, osteopathic, antirheumatic; vulnerary; auditory, tumour growth, retinal disorder; sports-related joint problem, articular carrilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 XP, Botstein D, Desnoyers L, Eaton DL;
E E, Fong S, Gao W, Gerber H, Gerritsen ME;
PJ, Grimaldi JC, Gurney AL, Hillan KJ;
Napler WA, Pan J, Paoni NF, Roy MA, Shelton I
Williams PM, Wood WI;
                                                                                                                                                                                                                                                                                                       / Match 2.9%; Score 12.4; DB 1; Length 18; Local Similarity 92.9%; Pred. No. 4.3e+02; nes 13; Conservative 0; Mismatches 1; Indels 0;
                                                                                                                                                                                                                                                                          Sequence 18 BP; 2 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
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22-DEC-1998; 98UG-0113296F.
08-MAR-1999; 99WG-US000106.
12-APR-1999; 99US-00284291.
25-AUC-1999; 99US-00284291.
18-FEB-2000; 2000WG-US004341.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human PRO 274 PCR primer #4.
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Ferrara N, Filvaroff E, F
Goddard A, Godowski PJ,
Kljavin IJ, Kuo SS, Napie
Stewart TA, Tumas D, Will
                                                                                                                                                                                                                                                                                                                                                                                           215 GAACTCGGTGGCGG 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADD72831 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                          18 GAACTCCGTGGCGG 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-JAN-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4PI; 2003-875643/81.
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                                                                                                                                                                                                                                                                                                               Query Match
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ADD72831/c
ID ADD7283
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19-OCT-2001; 2001US-00164929

Page 364

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Example 4; SEQ ID NO 14; 453pp; English.
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The invention relates to an isolated PRO polypeptide (secreted or transmembrane protein) having at least 60% amino acid sequence identity con an amino acid sequence chosen from 94 fully defined sequences as given in the specification (including PRO lacking its associated signal peptide). Also extracabilizar domain with or without its associated signal peptide). Also included are nucleic acids encoding the PRO proteins mentioned above, a vector comprising a PRO nucleic acid). A host cell comprising the vector and producing PRO a chimaeric molecule comprising the vector and producing PRO a chimaeric molecule comprising the vector and producing PRO a chimaeric molecule comprising the vector and producing PRO a chimaeric molecule comprising the vector and producing PRO a chimaeric molecule comprising the vector and producing PRO a chimaeric molecule comprising the vector and propertide is useful for detecting PRO159 polypeptide is useful for detecting PRO159 polypeptide; such propertide is useful for detecting PRO159 polypeptide; prof. PRO159 polypeptide is useful for detecting PRO159 polypeptide is useful for linking a bioactive molecule to a cell expressing PRO137 polypeptide; The bioactive molecule is the toxin, radiolabel, or an antibody. The bioactive molecule to a cell expressing PRO139 polypeptide is useful for linking a bioactive molecule is the toxin, radiolabel, or antibody propertide or acid acciding propertide is useful for linking a bioactive molecule to a cell expressing PRO159 polypeptide is useful for modulating at least one biological activity of the cell expressing PRO199 polypeptide or anti-PRO499 polypeptide is useful for modulating at least one biological activity of the cell expressing PRO199 polypeptide or anti-PRO499 polypepti

Sequence 18 BP; 2 A; 8 C; 4 G; 4 T; 0 U; 0 Other;

0; Gaps Query Match 2.9%; Score 12.4; DB 1; Length 18; Best Local Similarity 92.9%; Pred. No. 4.3e+02; Matches 13; Conservative 0; Mismatches 1; Indels

215 GAACTCGGTGGCGG 228 18 GAACTCCGTGGCGG 5 ò 엄

ADD72189,

RESULT 710

ADD72189 standard; DNA; 18 BP.

ADD72189;

29-JAN-2004 (first entry)

Human PRO 274 PCR primer #4.

Human, ss, PCR; secreted protein; transmembrane protein, PRO, cytostatic, ophthalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth; retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.

Homo sapiens.

US2003194781-A1

16-OCT-2003.

22-MAR-2001; 2001WO-US009552 25-MAY-2001; 2001WO-US017092 01-JUN-2001; 2001WO-US017800 09-JUL-2001; 2001WO-US02173 30-JUL-2001; 2001US-0091858 17-MAY-2000; 2 22-MAY-2000; 2 30-MAY-2000; 2 02-MAR-2000;

(GETH) GENENTECH INC.

Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL; Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME; Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ; Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL; Stewart TA, Tumas D, Williams PM, Wood WI;

WPI; 2003-852598/79.

New secreted and transmembrane PRO nucleic acids and polypeptides, useful for stimulating the release of tumor necrosis factor alpha from human blood and stimulating the proliferation of differentiation of chondrocyte

Example 4; SEQ ID NO 14; 462pp; English.

The invention relates to an isolated PRO polypeptide (secreted or transmembrane protein) having at least 80% amino acid sequence identity on an amino acid sequence chosen from 94 fully defined sequence as given in the specification (including PRO lacking its associated signal peptide, a PRO extracellular domain with or without its associated signal peptide, a Nso included are nucleic acids encoding the PRO proteins mentioned above, a vector comprising a PRO nucleic acids encoding the PRO proteins comprising the vector and producing PRO, a chimaeric molecule comprising antibody. PRO317 polypeptide is useful for detecting a PRO4993 polypeptide.

Similarly, PRO4993 polypeptide is useful for detecting PRO337

PRO725, PRO7020. PRO7039 polypeptide is useful for detecting
PRO155, PRO7030. PRO1559 polypeptide is useful for detecting
PRO155, PRO700 or PRO739. PRO4993 polypeptide is useful for linking a
bioactive molecule to a cell expressing PRO337 polypeptide. The bioactive
molecule is the toxin, radiolabel, or matibody. The bioactive molecule
causes death of the cell. PRO337 polypeptide is useful for linking a
bioactive molecule to a cell expressing PRO4993 polypeptide. PRO725,
PRO700 or PRO739 polypeptide are useful for linking a bioactive molecule
to a cell expressing PRO1559 polypeptide; and PRO1559 polypeptide is
useful for linking a bioactive molecule to a cell expressing PRO337

PRO700 or PRO739 polypeptide. PRO4933 polypeptide is
polypeptide is useful for modulating at least one biological activity of
the cell expressing PRO337 polypeptide, where the cell is killed. PRO337

polypeptide or anti-PRO4933 polypeptide is useful for modulating the piological activity of the cell expressing PRO355,

RO700 or PRO739 polypeptide or an anti-PRO4993 polypeptide; PRO700 or PRO725,

RO700 or PRO739 polypeptide or an anti-PRO4993 polypeptide; PRO700 or PRO725,

PRO700 or PRO739 polypeptide or an anti-PRO4993 polypeptide; DRO337

polypeptide; and PRO1559 polypeptide or anti-PRO525, anti-PRO700 or antiCC polypeptide; and PRO1559 polypeptide or anti-PRO705, anti-PRO700 or antiCC polypeptides are useful for inhibiting tumour growth, retinal disorders,

CC polypeptides are useful for inhibiting tumour growth, retinal disorders,

CC mammals. The present sequence is a PCR primer used to isolate nucleic acid encoding a PRO protein.

8899999999999999999

Sequence 18 BP; 2 A; 8 C; 4 G; 4 T; 0 U; 0 Other;

0; Gaps Score 12.4; DB 1; Length 18; Pred. No. 4.3e+02; 0; Mismatches 1; Indels Query Match
2.9%; Soc
Best Local Similarity 92.9%; Pre
Matches 13; Conservative 0;

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ADE16840 standard; DNA; 18 BP ADE16840,

ADE16840;

29-JAN-2004 (first entry)

Human PRO 274 PCR primer #4.

Human; ss; PCR; secreted protein; transmembrane protein; PRO; cytostatic; obthchalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth, retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.

Homo sapiens.

US2003203433-A1.

30-OCT-2003.

18-OCT-2001; 2001US-00145016

98US-0084414P. 98US-0113296P. 99WO-US000106. 99WO-US005028. 99US-00284291. 99US-00380138. 06-MAY-1998; 22-DEC-1998; 05-JAN-1999; 08-MAR-1999; 12-APR-1999;

(GETH) GENENTECH INC

25-AUG-1999; 99US-00380138. 18-FEB-2000; 2000WO-US004341. 30-JUL-2001; 2001US-00918585.

KP, Botstein D, Desnoyers L, Eaton DL;
E, Fong S, Gao W, Gerber H, Gerritsen ME;
PJ, Grimaldi JC, Gurney AL, Hillan KJ;
Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL;
Williams PM, Wood WI; Ferrara N, Filvaroff E, F Goddard A, Godowski PJ, G: Kljavin IJ, Kuo SS, Napiez Stewart TA, Tumas n

WPI; 2003-875640/81.

New genes, and its encoded secreted and transmembrane polypeptides, useful for treating e.g. lung or breast tumors, osteoarthritis, rheumatoid arthritis, obesity, diabetes, hyperinsulinemia, hypoinsulinemia or wounds.

Example 4; SEQ ID NO 14; 459pp; English.

The invention relates to an isolated PRO polypeptide (secreted or transmembrane protein) having at least 80% amino acid sequence identity to an amino acid sequence chosen from 94 fully defined sequences as given in the specification (including PRO lacking its associated signal peptide, a PRO extracellular domain with or without its associated signal peptide, a PRO extracellular domain with or without its associated signal peptide, a PRO extracellular domain with or without its associated signal peptide. Also included are nucleic acids encoding the PRO proteins mantioned above, a vector comprising a PRO nucleic acid), a host cell comprising the vector and producing PRO, a chimaeric molecule comprising to the vector and producing PRO, a chimaeric molecule comprising the vector and producing PRO, a chimaeric molecule comprising the vector and producing PRO, a chimaeric molecule comprising a sample suspected of containing PRO, appendix a sequence. Similarly, PRO, 4993 PRO, 4994 PRO, 4993 PRO, 4993 PRO, 4994 PRO, 49

Seguence 18 BP; 2 A; 8 C; 4 G; 4 T; 0 U; 0 Other;

acid encoding a PRO protein.

...allati 2.9%; Score 12.4; DB 1; Length 18; 92.9%; Pred. No. 4.3e+02; tive 0; Mismatches 1; Indels Local Similarity 92.9 Query Match Best Loca Matches

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ADE48348 standard; DNA; 18 29-JAN-2004 (first entry) ADE48348; ADE48348/ EXHXEXE

RESULT 712

Human PRO 274 PCR primer #4

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Human; ss; PCR; secreted protein; transmembrane protein; PRO; cytostatic; ophthalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth; retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.
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Homo sapiens.

US2003104536-A1

05-JUN-2003

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07-OCT-1998; 98WO-US021141.

20-NOV-1998; 98WO-US024655.

80-MAR.1999; 99WO-US00106.

80-MAR.1999; 99WO-US00106.

14-MAY.1999; 99WO-US005280.

14-MAY.1999; 99WO-US012252.

20-NOV-1999; 99WO-US012252.

20-DEC-1999; 99WO-US012252.

20-DEC-1999; 99WO-US01243.

20-DEC-1999; 99WO-US01243.

30-DEC-1999; 99WO-US01243.

30-MAR-2000; 2000WO-US00356.

30-MAR-2000; 2000WO-US00553.

30-MAR-2000; 2000WO-US014042.

30-MAR-2000; 2000WO-US014042.

30-MAR-2000; 2000WO-US014042.

30-MAR-2000; 2000WO-US014042.

30-MAR-2000; 2000WO-US014042.

30-MAR-2000; 2000WO-US014042.

30-MAR-2001; 2000WO-US013695.

30-MAR-2001; 2001WO-US013695.

30-MAR-2001; 2001WO-US013695.

30-MAR-2001; 2001WO-US013695.

30-MAR-2001; 2001WO-US019692.

30-MAR-2001; 2001WO-US01165.

30-MAR-2001; 2001WO-US01165.
19-OCT-2001; 2001US-00166709
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(GETH) GENENTECH INC

Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
Perrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ;
Kljavin IG, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton |
Stewart TA, Tumas D, Williams PM, Wood WI;

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WPI; 2004-008994/01.

New isolated nucleic acid encoding a PRO polypeptide, e.g. PRO4993 or PRO337, useful in molecular biology, chromosome and gene mapping, in generating antisense RNA and DNA, and in gene therapy.

Example 4; SEQ ID NO 14; 460pp; English.

The invention relates to an isolated PRO polypeptide (secreted or transmembrane protein) having at least 80% amino acid sequence identity

to an amino acid sequence chosen from 94 fully defined sequences as given competition (including PRO Jacking its associated signal peptide). Also including PRO Jacking its associated signal peptide. Also including are nucleic acids, and an anti-record mentioned above, a vector comprising a PRO nucleic acids, a host cell comprising the vector and producing PRO, a chimaeric molecule comprising the vector and producing PRO, a chimaeric molecule comprising the vector and producing PRO, a chimaeric molecule comprising the vector and producing PRO, a chimaeric molecule comprising the sample suspected of containing PRO4993 polypeptide is useful for detecting PRO370 polypeptide is useful for linking a productive molecule to a cell expressing PRO4993 polypeptide is useful for linking a productive molecule to a cell expressing PRO4993 polypeptide is useful for linking a bloactive molecule to a cell expressing PRO4993 polypeptide is useful for linking a causes death of the cell. PRO379 polypeptide is useful for linking a bloactive molecule to a cell expressing PRO4993 polypeptide is useful for linking a bloactive molecule to a cell expressing PRO4993 polypeptide is useful for linking a bloactive molecule to a cell expressing PRO4993 polypeptide or anti-PRO379.

C PRO700 or PRO739 polypeptide are useful for linking a bloactive molecule to a cell expressing PRO4993 polypeptide or anti-PRO379 colypeptide is useful for modulating at least one biological activity of the cell expressing PRO4993 polypeptide or anti-PRO370 or PRO799 polypeptide or an anti-PRO370 anti-PRO370 or PRO799 polypeptide or anti-PRO370 anti-PRO370 or PRO799 polypeptide or anti-PRO370 anti-PRO370 or PRO799 polypeptide or anti

Sequence 18 BP; 2 A; 8 C; 4 G; 4 T; 0 U; 0 Other;

.; 0 Query Match
2.9%; Score 12.4; DB 1; Length 18;
Best Local Similarity 92.9%; Pred. No. 4.3e+02;
Matches 13; Conservative 0; Mismatches 1; Indels

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215 GAACTCGGTGGCGG 228 18 GAACTCCGTGGCGG ò g

Human PRO 274 PCR primer #4.

Human, 88; PCR, secreted protein, transmembrane protein, PRO, cytostatic, ophthalmological; antiarthritic, osteopathic, antirheumatic; vulnerary; auditory, tumour growth, retinal disorder; sports-related joint problem, articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.

16-OCT-2001; 2001US-00978375

97US-0062250P.

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PR 13-NOV-1997; 97US-0065311P.
PR 21-NOV-1998; 98US-0077641P.
PR 11-NAR-1998; 98US-0077641P.
PR 11-NAR-1998; 98US-0077641P.
PR 11-NAR-1998; 98US-0077791P.
PR 12-MAR-1998; 98US-0077791P.
PR 20-MAR-1998; 98US-007791P.
PR 20-MAR-1998; 98US-0077891D.
PR 20-MAR-1998; 98US-0078930P.
PR 20-MAR-1998; 98US-0078930P.
PR 20-MAR-1998; 98US-0079863P.
PR 20-MAR-1998; 98US-0079863P.
PR 20-MAR-1998; 98US-0079863P.
PR 20-MAR-1998; 98US-0079863P.
PR 21-MAR-1998; 98US-0079863P.
PR 27-MAR-1998; 98US-0079862P.
PR 27-MAR-1998; 98US-008014P.
PR 27-MAR-1998; 98US-008014P.
PR 27-MAR-1998; 98US-008014P.
PR 27-MAR-1998; 98US-008014P.
PR 27-MAR-1998; 98US-0080132P.
PR 27-MAR-1998; 98US-0080133P.
PR 27-MAR-1998; 98US-0080133P.
PR 27-MAR-1998; 98US-0080133P.
PR 27-MAR-1998; 98US-0080133P.
PR 27-MAR-1998; 98US-0080139P.
PR 27-MAR-1998; 98US-0080139P.
PR 27-MAR-1998; 98US-008013P.
PR 27-MAR-1998; 98US-0080195P.
PR 27-MAR-19
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98US-0085580P 98US-0085680P 98US-0085680P 98US-0085680P 98US-0085680P 98US-0085610P 98US-0086610P 98US-00866114P 98US-00866114P 98US-00866114P 98US-00866114P 98US-00866114P 98US-00866114P 98US-00866114P 98US-0086611P 98US-0086611P 98US-008661P 98US-0103081 98US-01

15-MAY-1998;
15-MAY-1998;
15-MAY-1998;
15-MAY-1998;
15-MAY-1998;
15-MAY-1998;
12-MAY-1998;
12-MAY-1998;
12-MAY-1998;
12-MAY-1998;
12-MAY-1998;
12-MAY-1998;
12-MAY-1998;
12-MAY-1998;
12-MAY-1998;
13-MAY-1998;
14-MAY-1999;
16-UUL-1999;
16-UUL-1999;
17-MAR-1999;
18-MAR-1999;
18-MA

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This oligonuclectide is one of 13 sequences which hybridise to nucleotide sequences coding for recombinant tuberculosis antigenic polypeptides. When used as probes they could differentiate M. tuberculosis from other bacterial strains. When used as primers, the oligonuclectides amplify specific mycobacterial sequences (e.g. nucleotides 1-1358 of the BCG alpha-antigen). Amplified sequences are then detected using one of the complement of one of the other is primer site are claimed which comprise primer A(ii) with the complement of one of the other 13 primers. See also AAQ11081-3, AAQ11086, AAQ11086-90, AAQ11101-8, AAR11297-R11304. (Updated on 09-JAN-2003 to add missing OS field.) (Updated on 25-MAR-2003 to correct PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The sequence is that of hybridisation probe BGIA which was used as part of a method of simultaneously sequencing nucleic acids. (Updated on 25-MAR-2003 to correct PN field.)
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2.9%; Score 12.4; DB 1; Length 19;
Best Local Similarity 92.9%; Pred. No. 4.88+02;
Matches 13; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 1 A; 6 C; 8 G; 4 T; 0 U; 0 Other;
                    Claim 23; Page 66; 134pp; English.
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(first entry)
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Matches 13, Conservative
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15-JUN-1994
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AAQ54140/c
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Pred. No. 4.3e+02;
0; Mismatches 1;
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30-JUL-2001; 2001US-00918585
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(revised)
(first entry)
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Best Local Similarity 92.9
Matches 13, Conservative
                                             ASHKENAZI A J.
BAKER K P.
BOUSTEIN D.
BESNOYERS L.
EATON D L.
FERRARA N.
FILVRAROFF E.
FONG S.
                                                                                                                                                                                                                                                        GERBER H.
GERRITSEN M E.
GODDARD A.
GODOWSKI P J.
GIRMALDI J C.
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WILLIAMS P M.
WOOD W I.
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PAONI N F.
ROY M A.
SHELTON D L.
STEWART T A.
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KUO S S.
NAPIER M A.
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HILLAN K J.
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30-MAY-1991
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PANJ/)
PAON/)
ROYM/)
                                             (ASHK/)
(BOAE/)
(BOAE/)
(DESN/)
(FEAR/)
(FILV/)
(FONG/)
(GERB/)
(GERB/
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AAQ11087/c
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19 GGTGGCGGCCACAT

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The invention relates to new immortalised cell lines derived from preadipocytes containing an immortalising fragment of a viral oncogene. The immortalised adipocytes are used to identify substances able to regulate lipolysis and/or thermogenesis (potential therapeutic agents for treating diabetes and obesity). The cell lines have the advantage that they can be maintained in long term culture (contrast primary cultures of adipocytes) without loss of characteristic markers or ability to differentiate. The immortalised pre-adipocytes differentiate into mature adipocytes what placed in a medium containing insulin and dexamethasone. The primers AAT43089-19 are used to amplify marker genes to verify differentiation of the pre-adipocytes into mature adipocytes primers AAT43016-7 were used to amplify a second of the gene encoding a hormone sensitive lipase, a marker for mature "brown" adipocytes
                                                                                                                                                  Immortalised cell line; pre-adipocyte; viral oncogene; lipolysis; marker; thermogenesis; diabetees; obesity; cell culture; differentiation; mature; medium; insulin; dexamethasone; primer; PCR; polymerase chain reaction; amplification; hormone sensitive lipase; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immortalised pre-adipocytes contg viral oncogene fragment - useful for identifying cpds that regulate lipolysis and thermogenesis, as lipolytic agents and models for studying adipocyte processes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 12.4; DB 1; Length 19;
Pred. No. 4.8e+02;
0; Mismatches 1; Indels
                                                                                                                    Antisense primer to amplify hormone sensitive lipase gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP, 1 A, 9 C, 3 G, 6 T, 0 U, 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 17; 52pp; French.
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              AAT43117 standard; DNA; 19 BP.
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92.9%;
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Best Local Similarity 92.3.
Best Local Similarity 92.3.
Conservative
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                                                                                   05-SEP-1997
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                                                                                                                                                                                                                                                                                                             31-OCT-1996.
                                                                                                                                                                                                                                         Synthetic.
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                                                AAT43117;
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AAT16004
AAT4311
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Gaps ö

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AATIS996-16007 are allele-specific primers that were used to discriminate each of six HLA alleles that had been serologically typed in a LB33 cell line (a mealanoma cell line). Amplification refractory mutation system was used, which relies on a perfect nucleotide match at the 3' end of primers to ensure specificity of DNA amplifications. It was suggested that A24-B13.0c%, and A2B44-0ch consititute two HLA class I haplotypes of patient LB33, and that reduced expression of these haplotypes probably accounted for loss of antigen expression by immunoselected tumour cells. The invention concerns a nucleic acid (AAT08972) which encodes a tumour rejection antigen, which can be used in determining cancerous conditions in patients of tissue type HLA-B44
                                                                                                                                                                                                                                                                                                                                             Nucleic acid encoding tumour rejection antigen precursor - useful in assay for determining cancerous condition in patient of e.g. tissue type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                              tumour rejection antigen; detection; cancer; tissue type; HLA-B44; human leukocyte antigen; immunogenic; primer; PCR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tumour rejection antigen precursor; TRAP; HLA-Cw6; HLA-B44;
human leukocyte antigen B44; cytotoxic T lymphocyte; cancer;
therapy; diagnosis; vaccine; primer; PCR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
2.9%; Score 12.4; DB 1; Length 19;
Best Local Similarity 92.9%; Pred. No. 4.88+02;
Matches 13; Conservative 0; Mismatches 1; Indels
5' allele-specific primer for HLA-Cw6 amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 BP; 6 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HLA-Cw6 allele-specific 5' PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                              Example 9; Page 13; 44pp; English.
                                                                                                                                                                                                                                                    (LUDW-) LUDWIG INST CANCER RES.
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95US-00373636.
                                                                                                                                                                      95WO-US006852.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7 GAGTGAAACTGCGG 20
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(first entry)
                                                                                                                                                                                                                                                                                  Boon-Falleur T, Coulie P;
                                                                                                                                                                                                                                                                                                                   WPI; 1996-049316/05.
                                                                                                                                                                        31-MAX-1995;
                                                                                                                                                                                                      03-JUN-1994;
17-JAN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-MAR-2003
25-FBB-1998
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                                                                                                           W09533855-A1
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                                                                                                                                           14~DEC-1995
                                                                               Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                HLA-B44
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97WO-US001915.

96US-00602506.

Peyman A;

Cossum PA, Rando RF,

GR,

Revankar

96US-0015752P.

17-APR-1996;

(ARON-) ARONEX PHARM INC. Chaudhary N, Rao T, Uhlmann E; WPI; 1997-526457/48. for

Anti:sense oligonucleotide(s) inhibiting VEGF expression - used treating diseases characterised by vascularisation and vascular permeability, e.g. diabetic retinopathy.

Claim 40; Page 43; 64pp; English.

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05-FEB-1997;
 20-FEB-1996;
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2 Oligonucleotides (AAT79214 and AAT79215) respectively comprise 5' and 3' primers for the specific PCR amplification of HLA-Cw6 sequences.

Melanoma LB33 cell lines have been serologically typed as HLA-A24, A28, B13, B44, Cw6, Cw7. Semi-quantitative conditions for DNA amplification by PCR were established to assess the expression of each of the 6 class I alleles by different LB33-MEL tumour cell clones. Primers (AAT79206-17) were designed to enable discrimination of each allele from the 5 others. The results suggest that A24-B13-Cw6 and A28-B44-Cw7 constitute 2 HLA class I haplotypes accounts for loss of antigen exression by immunoselected tumour cells. Claimed tumour rejection antigens (see AAW23038-43) presented by HLA-B44 molecules can be used in methods for the diagnosis and therapy of cellular abnormalities involving expression of a tumour rejection antigen precursor, such as cancer, especially melanoma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense oligonucleotide, cellular vascular endothelial growth factor, VGFF, vascular permeability factor; blood vessel formation; anglogenesis; vascular permeability induction; disease progression; increased anglogenesis; phosphorothioate; ss.
                                                                                                                             Tumour rejection antigens presented by human leukocyte antigen B44 molecules - useful to identify HLA-B44 positive cells for diagnosis and therapy of cellular abnormalities.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Gaps
                                        Van Der Bruggen P, Luescher I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 12.4; DB 1; Length 19;
Pred. No. 4.8e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antisense oligonucleotide which inhibits VEGF expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 6 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                        Coulie P, Boonfalleur T,
                                                                                                                                                                                                                        Example 9; Page 15; 74pp; English
(LUDW-) LUDWIG INST CANCER RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВЪ.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7 GAGTGAAACTGCGG 20
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Best Local Similarity کوری
الم 13; Conservative
                                                                                    WPI; 1997-435086/40.
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                                        Herman J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT92948;
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AAT92948
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Novel antisense oligonucleotides AAT92942-62 reduce cellular vascular endothelial growth factor (VEGF) production in cells. Inclusion of a C5-propyryl uridine, or a C5-propyryl cytidine nucleotide residue in the oligonucleotide sequence increases the duplex melting temperature by at least 5 degrees celcius. VEGF, also known as vascular permeability factor. Is necessary for the formation of blood vessels (anglogenesis) carbot factor induces vascular permeability, is chemotactic for monocytes and osteoblasts, and is a selective mitogen for endothalial cells. Anormally high concentrations of VEGF are associated with diseases characterised by a high degree of vascularisation or vascular permeability. Cells tracted with the antisense oligonucleotides at concentrations of less that 1 micromolar, produce no more than 90% of the VEGF that is produced by untreated cells. The antisense oligonucleotides can be used for slowing the progression of diseases associated with creatment of diabetic retinopathy, aggressive cancers, psoriasis, reatment of diabetic retinopathy, aggressive cancers, psoriasis, returnatoid arthritis and other inflammatory conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HLA-B24; HLA-B44; tumour rejection antigen precursor; TRAP; cancer; human; melanoma; diagnosis; therapy; polymerase chain reaction; PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Match 2.9%; Score 12.4; DB 1; Length 19; Local Similarity 85.7%; Pred. No. 4.8e+02; les 12; Conservative 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Coulie P, Van Der Bruggen P, Boonfalleur T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 5 A; 5 C; 6 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (LUDW-) LUDWIG INST CANCER RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HLA-Cw6 allele 5' PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAT62046 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            39 GAAGATGGCCACCA 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
(first entry)
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29-OCT-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
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Matches
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/*tag= a /note= "Phosphorothioate linkage between each residue; C5

-propynyl pyrimidines"

WO9739120-A2

23-OCT-1997

97WO-US006412

17-APR-1997;

Location/Qualifiers

/*tag=

Key modified_base Homo sapiens.

Synthetic

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Wed Apr 21 12:58:21 2004
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WPI; 1997-202614/18.

 $\rm HLA-B44~molecule$ binding peptide(s) - useful to identify $\rm HLA-B44~positive$ cells, and develop products for diagnosis and therapy of, e.g. cancer.

Example 9; Page 13; 55pp; English.

This 5' primer is specific for class I allele HIA-Cw6. Allele- specific primers (AATGO18-49) enable discrimination of each of the six class I alleles (HIA-A24, A28, B13, B44, Cw6 and Cw7) of melanoma patient LB33. DNA from different LB33-WEL tumour cell clones was subjected to PCR amplification. The results showed that A24-B13-Cw6 and A28-B44-Cw7 constitute two HIA class I haplotypes of patient LB33, and that reduced expression of these haplotypes probably accounts for loss of antigen expression by immunoselected tumour cells. HIA-B44 binding peptides (AAW1251-S6) can be used to identify HIA-B44 boiltive cells, and to develop products for the diagnosis and therapy of e.g. cancer, particularly melanoma. (Updated on 25-WAR-2003 to correct PI field.)

Sequence 19 BP; 6 A; 5 C; 6 G; 2 T; 0 U; 0 Other;

Gaps . 0 Ouery Match 2.9%; Score 12.4; DB 1; Length 19; Best Local Similarity 92.9%; Pred. No. 4.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels

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AAX59110 standard; DNA; 19 BP. AAX59110; RESULT 721 AAX59110/c ID AAX591

31-AUG-1999 (first entry)

Human nuclear receptor nNR5 PCR primer R5R4.

Nuclear receptor protein; nNR5; human; retina; eye disease; therapy; diagnosis; PCR; primer; ss.

Synthetic. Homo sapiens.

17-JUN-1999

98WO-US026422 11-DEC-1998;

12-DEC-1997;

(MERI) MERCK & CO INC.

Chen F;

WPI; 1999-385576/32.

DNA encoding human nuclear receptor nNR5.

Example 1; Page 32; 57pp; English.

This oligonuclectide comprises PCR primer R5R4, which was used with primer R5F3 (see AAX59109) to define the intron-exon boundary in a cDNA clone (see AAX5908) that had been isolated from a human retina cDNA library and which coded for a novel member of the nuclear receptor superfamily. An intronless clone (see AAX59095) was subsequently a morel member of the human nuclear factor superfamily. In the retina cDNA library. This encoded nR78 (see AAY06301), a novel member of the human nuclear factor superfamily. NRFs is expressed at high levels in the retina and may therefore play a role in eye

function. The invention also provides recombinant vectors and host cells, methods of screening for modulators of nNR5 activity, and production of antibodies against nNR5 88888

Sequence 19 BP; 2 A; 7 C; 8 G; 2 T; 0 U; 0 Other;

ô Gaps .; 0 Length 19; 1; Indels 2.9%; Score 12.4; DB 1; 92.9%; Pred. No. 4.8e+02; tive 0; Mismatches 1; Query Match Best Local Similarity 92.99 Matches 13; Conservative

ઠ 용 RESULT 722 AAZ87065/c

AAZ87065 standard; DNA; 19 BP.

AAZ87065;

(first entry) 16-MAY-2000

RBP-7 microsequencing primer for marker 5-143-84.

RBP-7; retinoblastoma binding protein-7; abnormal cell proliferation; diagnosis; therapy; cell differentiation; thyroid hyperplasia; psoriasis; benign prostate hypertrophy; cancer; sarcoma; neoplasm; leukaemia; lymphoma; biallelic marker; primer; ss.

Homo sapiens

WO200000607-A1.

06-JAN-2000.

99WO-IB001242. 30-JUN-1999; 98US-0091315P. 98US-0111909P. 30-JUN-1998; 10-DEC-1998;

(GEST) GENSET

Bougueleret L;

WPI; 2000-117170/10.

Novel nucleic acid and polymorphic markers used for diagnosis of diseases, especially those involving abnormal cell proliferation and differentiation.

Claim 15; Page 218; 223pp; English.

This sequence represents a microsequencing primer for a biallelic marker from the retinoblastoma binding protein-7 (RBP-7) genomic sequence (AAZ86567) of the invention. The RBP-7 coding sequence and regulatory caquesses are useful for the recombinant production of the protein and for expressing heterologous nucleic acids. Primers and probes derived from the RBP-7 nucleotide sequence (such as this sequence) are useful for DNA amplification and detection methods. RBP-7 biallelic markers (see AAZ86993-Z87034) are useful for diagnosis of disease related to alteration in the regulation or in the coding regions of the RBP-7 gene and for prognosis of an eventual treatment with therapeutic and especially agents acting on pathologies involving abnormal cell proliferation and/or differentiation, these include thyroid hyperplasis, psoriasis, benign prostate hypertrophy, cancers, including breast cancer, prostate cancer various leukaemias, and lymphomas. RBP-7 antibodies are useful as diagnostic agents

Seguence 19 BP; 3 A; 9 C; 3 G; 4 T; 0 U; 0 Other;

2.9%; Score 12.4; DB 1; Length 19;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention provides a database of human samples obtained from healthy individuals which can be used to identify polymorphic genetic markers. Data obtained for the database can be used to sort the samples by parameters such as age, sex and ethnicity. This is useful in linking markers with diseases, susceptibility to infection and drug responses. The present sequence was used in an assay to demonstrate the uses of the database of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Producing a database for identifying polymorphic genetic markers, comprises obtaining data relating to members of a healthy population and entering the information into a database.
                                                                                                                                                                                                                                                                  Database; polymorphism; SNP; human; genetic marker; disease; infection; drug response; ds.
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                                                                                                                                                                                                                                           Human lipoprotein lipase coding sequence fragment SEQ ID NO: 6.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 2.9%; Score 12.4; DB 1; Length 19; Best Local Similarity 92.9%; Pred. No. 4.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels
                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rodi C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 2 A; 9 C; 6 G; 2 T; 0 U; 0 Other;
Pred. No. 4.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ping Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Koester H, Van Den Boom D,
Jurinke C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 182; 304pp; English
                           0
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                                                                                                                                                                                                                                                                                                                                                                                                                               13-OCT-1999; 99US-0159176P.
10-ULL-2000; 2000US-0217551P.
10-ULL-2000; 2000US-0217658B.
19-SEP-2000; 2000US-00663968.
                                                                                                                                                                                                                                                                                                                                                                                                    13-OCT-2000; 2000WO-US028413
            92.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               238 GAGGCTGCTTCCCG 251
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABL53403 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          s caccircarcacce 18
                                                                                                                                                        AAH02312 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                        55 CAGAGGAGTCTCTG 68
                                                                                                                                                                                                                 (first entry)
                                                                               14 CAGAGGAGTCACTG 1
            Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQU-) SEQUENOM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-273865/28
                                                                                                                                                                                                                                                                                                                                              WO200127857-A2.
                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-MAY-2002
                                                                                                                                                                                                               12-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                          19-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABL53403;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Braun A,
                                                                                                                                                                                    AAH02312;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 724
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABL53403
                                                                                                                          RESULT
AAH02312
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The invention relates to a protein related to the hemagglutination or hemadsorption reaction and the gene thereof are provided to screen effective candidates in the diagnosis of fungal infection. The protein of the invention performs coagulation of foreign material. A phenoloxidase abenoication and non-self selectively. The phenoloxidase and the gene thereof induce cagulation and adsorption of a foreign material by using blood cell so as to block diffusion. The protein selectively removes fungi and bacteria invading a body, and is used in the diagnosis of pathogenic foreign material. A pro-phenoloxidase is used in detection of melanin forming represent: The current sequence represents haemagglutination or haemadsorption related DNA referred to as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               F2718; miz-1; forward PCR primer; Leishmania tarentolae; proto-oncogene;
8s; pIR-BgIII-forward; colony-PCR.
                                                                                                                                                                                                                                                                                  Protein related to hemagglutination or hemadsorption reaction and genethereof.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 12.4; DB 1; Length 19;
Pred. No. 4.8e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      F2718 (pIR-BgIII-forward) miz-1 proto-oncogene PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 5 A; 5 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                Park JJ;
                                                                                                                                                                                                                                Lee HS,
                                                                                                                                                                                                                                                                                                                                Disclosure; Page 7; 9pp; Korean.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAH24334 standard; DNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-NOV-2000; 2000WO-EP010794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99EP-00122222
                                                                                                                                                                            99KR-00026408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 2.9%;
Best Local Similarity 92.9%;
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          39 GAAGATGGCCACCA 52
                                                                                                                                                                                                       (SAMY-) SAMYANG GENEX CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 GAAGAGGGCCACCA 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                            WPI; 2001-472806/51
                                                                                                                                                                                                                                   Lee BR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200132896-A1
                                                                                           KR2001005404-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-NOV-1999;
                                                                                                                                                28-JUN-1999;
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                                                                                                                                                                            28-JUN-1999;
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                                                                                                                       LS-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                Hong SS,
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Haemagglutination; haemadsorption; fungal infection; phenoloxidase; ds.

Haemagglutination or haemadsorption related DNA SK-2.

WPI; 2001-316448/33.

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The present sequence represents PCR forward primer-F2718 (pIR-BgIII-forward), used to identify clones of B. coli TG1 which had been transformed with the pIR plasmid containing the miz-1 sequence insert in the right orientation. The present sequence was used to identify these clones using colony-PCR. This was part of an experiment of the invention to express miz-1 in Leishmania tarentolae. The invention comprises an expression and delivery system for the production of recombinant protein with cultivated non-pathogenic Kinetoplastidae parasites. The invention is used to express heterologous proteins or to deliver heterologous proteins into plant or animal cells, unlike prior art recombinant protein expression in Kinetoplastidae, this invention uses non-pathogenic species which do not carry the associated health risks, does not require expensive, uncommon media, and has a relatively high growth rate
A new recombinant protein expression system using non pathogenic
Kinetoplastidae type host cells such as Leishmania tarentolae allows
large scale production on inexpensive media.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Match 2.9%; Score 12.4; DB 1; Length 19; Local Similarity 92.9%; Pred. No. 4.8e+02; es 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 2 A; 7 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                           Example 1; Page 15; 37pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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Matches
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0; Gaps
                                                                                                                                                                                                                                         Chimeric oligonucleotide Kan4021C fragment #2.
                                 255 TCGGCCACGGTGCA 268
                                                                                                                                            ABQ79482 standard; DNA; 19
                                                                                                                                                                                                           15-NOV-2002 (first entry)
                                                              15 TCGACCACGGTGCA 2
                                                                                                                                                                              ABQ79482;
                                                                                                                 RESULT 726
                                   ò
                                                                g
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Mutation; genetic repair; point mutation; frameshift mutation; plant; plastid; chloroplast; repair; DNA recombination; ss. 07-JAN-2002; 2002WO-US000338. 05-JAN-2001; 2001US-0260076P. WO200259380-A2. 01-AUG-2002. Synthetic.

Kmiec EB; May GD,

(ROBE-) ROBERTS NOBLE FOUND INC SAMMUEL

WPI; 2002-599808/64.

Modifying a target site of a plastid gene-of-interest, useful for plant genetic repair, comprises reacting chimeric RNA/DNA oligonucleotides or modified DNA oligonucleotides in conjunction with a cell-free chloroplast lysate.

Example 1; Fig 2; 28pp; English.

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The invention relates to an in vivo or in vitro cell-free method for genetic repair of mutations in plastid genes. The method involves confecting a plastid which contains a specific point or frameshift mutation of interest, a chimeric RNA/DNA oligomucleotide or a modified single confection or contain the genetic code (for correcting the gene mutation, and a chloroplast extract taken from the plant of interest. The method of the invention is useful for plant of energis, and facilitates the direct comparison between plant unclear comparison between plant mutera and organelle DNA repair pathways. The cell-free assay may be used in clucidating plastid DNA recombination and repair pathways in plant cells as well as the identification and characterisation of proteins involved in the process. The current sequence represents a fragment of the chimeric oligomucleotide Kan4021C. This sequence is used in an example commert of in the process. The current sequence represents a fragment of the chimeric oligomucleotide Kan4021C. This sequence is used in an example contained in plasmid pksm4021 is converted in order to restore canamycin resistance activity. The chimeric oligomucleotide fragment conversion has occurred
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Sequence 19 BP; 4 A; 5 C; 6 G; 4 T; 0 U; 0 Other;

; / Match 2.9%; Score 12.4; DB 1; Length 19; Local Similarity 92.9%; Pred. No. 4.8e+02; nee 13; Conservative 0; Mismatches 1; Indels 155 ÇGGÇTTÇGAÇTÇGG 168 4 cecraceaciese 17 Query Match Best Local Si Matches 13; ઠે 셤

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ВР. ADA25739 standard; RNA; 19 ADA25739; RESULT 727 ADA25739/

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20-NOV-2003 (first entry)

short interfering nucleic acid; siNA; nuclear factor kappa B; NF-kappaB; RNA interference; vasotropic; noctropic; antiparkinsonian; neuroprotective; cytostatic; antiinflammatory; antiallergic; virucide; anti-HUV; immunosuppressive; anticonvulsant; nephrotropic; gene therapy; modulation; inhibition; restenosis; central nervous system lesion; Alzheimer's disease; Parkinson's disease; Huntington's disease; epilepsy; polycystic kidney disease; inflammatory disease; allergic disease; viral infection; HIV; autoimmune disease; transplant rejection; ribozyme; human; v-rel reticuloendotheliosis viral oncogene homologue A; REL-A; nuclear factor; ss.

20-FEB-2002; 2002US-0358580P. 11-MAR-2002; 2002US-0363124P. 06-UUN-2002; 2002US-0386782P. 29-AUG-2002; 2002US-0406784P. 05-SEP-2002; 2002US-0408378P. 09-SEP-2002; 2002US-0409293P. 20-FEB-2003; 2003WO-US004951 LS-JAN-2003; 2003US-0440129P WO2003070970-A2. Homo sapiens. 28-AUG-2003.

Beigelman L; Mcswiggen J,

Human REL-A short interfering nucleic acid SEQ ID NO:87.

Synthetic.

(RIBO-) RIBOZYME PHARM INC.

(JENA-) JENA BIOSCIENCE GMBH. Alexandrov K,

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WPI; 2003-689788/65.
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New short interfering nucleic acid downregulates expression of the NP-kappaB gene useful e.g. for treatment and diagnosis of cancer and
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Example 3; Page 129; 149pp; English

29-AUG-2002; 2002US-0406784P. 05-SEP-2002; 2002US-0408378P. 09-SEP-2002; 2002US-0409293P. 15-JAN-2003; 2003US-0440129P.

(RIBO-) RIBOZYME PHARM INC.

Mcswiggen J, Beigelman L; WPI; 2003-689788/65.

20-FEB-2002; 2002US-0358580P. 11-MAR-2002; 2002US-0363124P. 06-JUN-2002; 2002US-0386782P. 20-FEB-2003; 2003WO-US004951

28-AUG-2003

The present invention describes a short interfering nucleic acid (siNA)

that downregulates expression of a nuclear factor kappa B (NF-kappaB)

that downregulates expression of a nuclear factor kappa B (NF-kappaB)

control by RNA interference. Also described: (1) Kits for in vitro or in

vivo delivery of siNA; (2) conjugates and/or complexes of siNA; and (3)

control control of expression or complexes of siNA; antinflammatory,

antiallargic, virucide, and can be used in gene therapy, and for the

nephrotropic activities, and can be used in gene therapy, and for the

nephrotropic activities, and can be used in gene therapy, and for the

condulation (inhibition) of expression or activity of NF kappaB by RNA

interference (siNA target mRNA, pre-RNA and/or RNA templates). The siNA

concers that transplants for treating restenosis and central nervous system

colls, tissue explants for treating restenosis and central nervous system

colls, tissue explants for treating restenosis and central nervous system

colls, tissue explants for treating restenosis and central nervous system

collessons and injuries (Alataimer's, Parkinson's or Huntington's diseases,

collessons and injuries (Alataimer's, Parkinson's or furting and polycystic kidney

disease, inflammatory and/or allergid diseases, viral infections

concers, other proliferative diseases (restenosis and polycystic kidney

disease), inflammatory and/or allergid diseases, viral infections

concers, other proliferative diseases and transplant rejection, and also

concers, other proliferative diseases and transplant rejection, and solor

concers, other proliferative diseases sudding gene function and gene

concers, other proliferative diseases sudding gene

conce

Sequence 19 BP; 1 A; 9 C; 8 G; 0 T; 1 U; 0 Other;

ö 0; Gaps Ouery Match
2.9%; Score 12.4; DB 1; Length 19;
Best Local Similarity 92.9%; Pred. No. 4.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels 13; Conservative

ADA26088;

short interfering nucleic acid; siNA, nuclear factor kappa B; NP-kappaB; RNA interference, vasctropic; nootropic; antiparkinsonian; neuroprotective, cytostatic; antinflammatory; antiallergic; virucide, anti-HUV; immunosuppressive; anticonvulsant; nephrotropic; gene therapy; modulation; inhibition; restenosis; central nervous system lesion; Alzheimer's disease; Parkinson's disease; Huntington's disease; epilepsy; dementia; amyotrophic lateral sclerosis; cancer; polycystic kidney disease; inflammatory disease; allergic disease; viral infection; HUV; autoimmune disease; transplant rejection; ribozyme; human; verel reticuloendotheliosis viral oncogene homologue A; REL-A; nuclear factor; ss.

domo sapiens. Synthetic

402003070970-A2

305 GAGCCCCGGGGACC 318 15 GAGCCCCGGGGCCC 2 셤 ò

Human REL-A short interfering nucleic acid SEQ ID NO:223. ADA26088 standard; RNA; 19 BP. 20-NOV-2003 (first entry) RESULT 72 ADA26088

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The present invention describes a short interfering nucleic acid (siNA) that downregulates expression of a nuclear factor kappa B (NF-kappaB) con the downregulates expression of a nuclear factor kappa B (NF-kappaB) con the factor should be the special by NAA interference. Also described: (1) kits for in vitro or in vivo delivery of siNA; (2) conjugates and/or complexes of siNA; and (3) confucing siNA. The siNAs have vasotropic, nootropic, notropic activities, and can be used in gene therapy, and for the nephrotropic activities, and can be used in gene therapy, and for the interference (siNA target mRNA, RNA splite variants, post-circulation (inhibition) of expression or activity of NF-kappas by RNA interference (siNA target mRNA, pre-RNA and/or RNA templates). The siNA captures con organisms, e.g. by ex vivo gene therapy, in cells, tissue explants or organisms, e.g. by ex vivo gene therapy, in cells, tissue explants for treating restencis and central nervous system and injudies (Alzheimer's, Parkinson's or Huntington's diseases, epilepsy, dementia or amyotrophic lateral sclerosis or for treating many cancers, other proliferative diseases (restencis and polycystic kidney cancers, other proliferative diseases (restencis and polycystic kidney (including HIV), autoimmune diseases and transplant rejection, and also for drug screening; diagnosis; target identification and validation; contineering; pharmacogenomics; studying gene function and gene contral nervents human v-rel reticulcendcheliosis viral oncogene homologue A (REL-A) sinA, which is used in the exemplification of the present envented in B-cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                               New short interfering nucleic acid downregulates expression of the NF-kappaB gene useful e.g. for treatment and diagnosis of cancer and inflammation.
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2.9%; Score 12.4; DB 1; Length 19;
Best Local Similarity 92.9%; Pred. No. 4.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3; Page 129; 149pp; English.
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ADD00605 standard; RNA; 19 BP. 5 GAGCCCCGGGGCCC 18 01-JAN-2004 (first entry) ADD00605; RESULT 729 g

HCV infection; replication; pathogenesis; virucide; vaccine; HCV coding region-derived 50% conserved RNA sequence 551.

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Polynucleotides encoding nuclear receptors, and the encoded proteins, useful as diagnostic agents, and for identification of agents that affect receptor activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to novel nuclear receptor ERR (oestrogen receptor-
related receptor) gamma 3 polynucleotides. The polynucleotides of the
invention may be useful for diagnosis of disorders caused by abnormal
nuclear receptor activity, particularly those related to abnormal
                                                                                                                                                                                                                     The invention relates to a novel isolated double stranded RNA oligonucleotide about 19 to about 25 ribonucleotide comprises the same equivalent. One strand of the oligonucleotide comprises the same nucleotide sequence as region of a hepatitis C virus (HCV) target RNA polynucleotide sequence required for hepatitis C virus infection, replication or pathogenesis in vitro or in vivo in a host cell. The oligonucleotide of the invention demonstrates virucide activity and may be useful for preparing a composition or vaccine for treating or preventing hepatitis C virus, as well as during gene therapy procedures. The current sequence is that of the HCV coding region-derived conserved RNA sequence of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nuclear receptor; ERR gamma 3; oestrogen receptor-related receptor; oestrogen receptor; ER; thyroid hormone; TR; human; ss; PCR; primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1; Indels 0;
                                                                                                                      New double stranded RNA oligonucleotide, useful for preparing composition for treating or preventing hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
2.9%; Score 12.4; DB 1; Length 19;
Best Local Similarity 92.9%; Pred. No. 4.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 4 A; 6 C; 2 G; 0 T; 7 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human ERR gamma 3-related PCR primer - SEQ ID 13.
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                                        Yang
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                                             Martinez A,
                                                                                                                                                                                          Disclosure; Page 90; 173pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6 GGAGTGAAACTGCG 19
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                                           Glass JI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGAGTGAAAATGCG
(ELIL ) LILLY & CO ELI.
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                                                                                      WPI; 2003-268345/26.
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                                             Ľu J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
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                                             Zhao G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 731
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a novel isolated double stranded RNA oligonucleotide about 19 to about 25 ribonucleotides in length or its equivalent. One strand of the oligonucleotide sin length or its nucleotide sequence as a region of a hepatitis C virus (HCV) target polynucleotide sequence required for hepatitis C virus infection, replication or pathogenesis in vitro or in vivo in a host cell. The oligonucleotide of the invention demonstrates virucide activity and may be useful for preparing a composition or vaccine for treating or preventing hepatitis C virus, as well as during gene therapy procedures. The current sequence is that of the HCV coding region-derived conserved RNA sequence of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                New double stranded RNA oligonucleotide, useful for preparing a composition for treating or preventing hepatitis C virus.
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2.9%; Score 12.4; DB 1; Length 19; 92.9%; Pred. No. 4.8e+02; ive 0; Mismatches 1; Indels

Local Similarity 92.9%; Les 13; Conservative

Query Match

Best Loc Matches

6 GGAGTGAAACTGCG 19

19 GGAGTGAAAATGCG 6

g

8

Sequence 19 BP; 3 A; 7 C; 2 G; 0 T; 7 U; 0 Other;

Martinez A, Yang Y;

Lu J, Glass JI,

Zhao G,

WPI; 2003-268345/26.

(ELIL) LILLY & CO ELI

16-AUG-2002; 2002WO-US021843. 17-AUG-2001; 2001US-0313076P. 20-DEC-2001; 2001US-0344116P. 01-FEB-2002; 2002US-0353750P.

Hepatitis C virus.

WO2003016572-A1.

27-FEB-2003

gene therapy; ds.

Disclosure; Page 90; 173pp; English.

infection; replication; pathogenesis; virucide; vaccine;

Hepatitis C virus.

WO2003016572-A1.

27-FEB-2003

gene therapy; ds

HQ.

16-AUG-2002; 2002WO-US021843. 17-AUG-2001; 2001US-0313076P. 20-DEC-2001; 2001US-0344116P. 01-FEB-2002; 2002US-0353750P.

HCV coding region-derived 50% conserved RNA sequence 552.

01-JAN-2004 (first entry)

ADD00606;

BP.

ADD00606 standard; RNA; 19

730

ADD00606, RESULT

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Hepatitis C virus; non-A non-B virus; HCV-Hc59; primers; probes; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a method of identifying a class I or II Human Leukocyte Antigen (HLA) genotype of a subject using hybridisation and amplification assay. The method is used for determining the HLA genotype of a subject. The present sequence represents a HLA class I allele
                                                                                                                                                                                                                                                                                                                                                                                                          Identifying class I or II Human Leukocyte Antigen genotypes using hybridization and amplification assays.
                                                                                    ss; primer; PCR; human; Human Leukocyte Antigen; HLA; genotype
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          . Match 2.9%; Score 12.4; DB 1; Length 19; Local Similarity 92.9%; Pred. No. 4.8e+02; les 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 6 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                  HLA class I allele specific primer #117
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 7; SEQ ID NO 119; 66pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HCV-Hc59 primer #795 (sense strand).
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90US-00616369.
91US-00748564.
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                                                                                                                                                                                                                       25-APR-2002; 2002US-00133779.
                                                                                                                                                                                                                                                     20-DEC-1999; 99US-0172768P.
20-DEC-2000; 2000US-00747391.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ22903 standard; DNA; 17
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                   (first entry)
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                                                                                                                                                                                                                                                                                                        (STEM-) STEMCYTE INC
                                                                                                                                                                                                                                                                                                                                          Tonai R;
                                                                                                                                                      US2003165884-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  specific primer
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21-AUG-1991;
                                                                                                                       Homo sapiens.
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                   29-JAN-2004
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07-JUL-1992
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                                                                                                                                                                                      04-SEP-2003.
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Matches
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oestrogen receptor (ER), ERR or thyroid hormone receptor (TR) activity. Furthermore, the polynuclectides and proteins may be useful for evaluating agents that affect the activity of nuclear receptors. The current sequence is that of the human ERR gamma 3-related PCR primer (ID 13) of the invention.
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                                                                                                                                     Length 19;
                                                                                                                                                                      1; Indels
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                                                                                                   Sequence 19 BP; 3 A; 7 C; 2 G; 7 T; 0 U; 0 Other;
                                                                                                                                   Query Match
2.9%; Score 12.4; DB 1;
Best Local Similarity 92.9%; Pred. No. 4.8e+02;
Matches 13; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                               HLA class I allele specific primer #1.
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                                                                                                                                                                                                                                                                                                                            ADE13385 standard; DNA; 19 BP.
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20-DEC-2000; 2000US-00747391
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                                                                                                                                                                                                                              6 TCCCTGCACTACGA 19
                                                                                                                                                                                                          63 TCTCTGCACTACGA 76
                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tonai R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             specific primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                 29-JAN-2004
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Best Local S
                                                                                                                                                                                                                                                                                                                                                              ADE13385;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 733
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ID ADE
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AC ADE
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Gaps

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(PHAR-) PHARMA
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One Hutch strain (HCV-H) of NANBV, designated the Hutch c59 isolate (HCV-HC59) was propagated through passage in animals and the entire viral genome was cloned and sequenced. Five microg of purified liver or plasma derived from HCV RNA was used per CDNA priming reaction. Specific nucleotide primers derived from published HCV sequences and spanning the entire reported genomic sequences were used to prime the reaction. Selected target sequences were amplified using a PCR-based approach using a variety of nucleotide primers. The nucleotide sequences of the primers subsequently isolated, randered blunt-ended and inserted into a pUC or pBluescript cloning vectors. (Updated on 25-MAR-2003 to correct PR field.) (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                     Deoxyribonucleic acid sequence encoding non-A, non-B hepatitis virus - obtd. Hutch C59 subgroup encoding polypeptide(s), useful as vaccines, and immuno reactive ABS for diagnosis of virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 1 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
Prince AM;
Nasofe MS,
                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 107; 225pp; English
Zebedee S, Inchauspe G,
                                                                                  WPI; 1992-096821/12
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Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                266
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                                                                              250 CGGGCTCGGCCACGGTG
                                                                                                                 g
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0; Gaps

Human mismatch repair pathway gene MSH2, primer 17209. 383/c AAQ92383 standard; DNA; 17 BP (first entry) 15-JAN-1996 AAQ92383; AAQ92383,

Mismatch repair; MSH2; primer; identification; defect; alteration; cancer; tumour; vaccine; 88. 93US-00154792. 93US-00163449. 94US-00259310. 94WO-US013385 Homo sapiens WO9514085-A2 7-NOV-1994; 7-NOV-1993; 26-MAY-1995

(DAND) DANA FARBER CANCER INST. (UTVE-) UNIV VERMONT & STATE AGRIC COLLEGE.

13-JUN-1994;

17-DEC-1993;

Fishel R, Reenan RA; Colodner RD,

WPI; 1995-200377/26.

Determining alteration in human mismatch repair pathways - used in the diagnosis, prognosis and therapy of cancers and in screening assays.

Claim 15; Page 186; 256pp; English

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AAQ92382-Q92400 and AAQ93890-Q93900 are oligonuclectide primers used to detect alterations in the human mismatch repair pathway gene MSH2. Defects or alterations in such a gene result in the accumulation of unstable repeated DNA sequences, a feature of a number of different cancers. The identification of a defect in the mismatch repair pathway can be diagnostic of a predisposition to cancer and prognostic for a particular mammalian cancer e.g colorectal, ovarian, endometrial (uterine), renal, bladder, skin, rectal and bowel. The nucleotide sequences and polypeptides of the hMSH2 gene may also be used for therapy and in vaccines
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Sequence 17 BP; 2 A; 10 C; 3 G; 2 T; 0 U; 0 Other;

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Query Match
2.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 82.4%; Pred. No. 4.2e+02;
Matches 14; Conservative 0; Mismatches 3; Indels
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319 GCGTGCTGGCGGCGGAC 335 17 dedrigerigedadedae ઠે 셤

AAX74482 standard; RNA; 17 BP. RESULT 736 AAX74482

AAX74482;

(first entry) 28-JUL-1999

Mouse flt-1 VEGF receptor hammerhead ribozyme substrate #10

Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; ss.

Mus sp.

WO9715662-A2. 01-MAY-1997. 96WO-US017480. 25-OCT-1996;

95US-0005974P. 96US-00584040. 26-OCT-1995; 11-JAN-1996;

(RIBO-) RIBOZYME PHARM INC. (CHIR) CHIRON CORP.

Pavco P, Mcswiggen J, Stinchcomb D, Escobedo J;

WPI; 1997-259017/23.

Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.

Claim 4; Page 155; 218pp; English.

The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention

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The present invention describes enzymatic nucleic acid molecules (NAMs) which specifically cleave RNA derived from an epidermal growth factor receptor (EGR-R) gene. AAV09721 to AAV096043 and AAV09999 to AAV099999 represent specifically claimed target sequence from human EGF-R. AAV098044 to AAV98866 and AAV98867 to V9978 represent hammerhead ribozymes and hairpin ribozymes respectively for human EGF-R. The NAMs are useful for cleaving EGF-R. RNA in the treatment of a condition associated with EGFR expression levels e.g. to inhibit cell proliferation in the prevention or treatment of cancers. The NAMs can also be used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of EGF-R RNA in a cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BDIXymatic nucleic acids - which cleave RNA derived from an epidermal
growth factor receptor, useful for inhibiting cell proliferation and for
treating cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human, epidermal growth factor receptor; EGFR; EGF-R; target sequence; hammerhead ribozyme; hairpin ribozyme; inhibition; cell proliferation; cancer; genetic drift; detection; mutation; ss.
                                                                                                                    Human, epidermal growth factor receptor; EGFR; EGF-R; target sequence;
hammerhead ribozyme, hairpin ribozyme; inhibition; cell proliferation;
cancer; genetic drift; detection; mutation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human EGF-R target sequence nucleotide position 4356.
                                                                                Human EGF-R target sequence nucleotide position 4357.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 5; Page 79; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mcswiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      390 GGCGCCAAGAAGGTCTT 406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV97773 standard; RNA; 17 BP.
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97US-00985162.
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                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                     14-JAN-1998;
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                                            17-MAR-1999
                                                                                                                                                                                                                                                                                               06-AUG-1998
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    AAV97774;
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AAV97773/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Treatment of airway diseases such as asthma - by topically applying adenosine-free antisense oligo:nucleotide to airway epithelium of subject.
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llarity 82.4%; Pred. No. 4.2e+02;
Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                          Asthma; airway epithelium; adenosine free; cystic fibrosis; chronic obstructive pulmonary disease; bronchitis; ss.
                                      Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 58.8%; Pred. No. 4.2e+02; Matches 10; Conservative 4; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                    Endothelial nitric oxide antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 0 A; 7 C; 6 G; 4 T; 0 U; 0 Other;
  Sequence 17 BP; 0 A; 5 C; 5 G; 0 T; 7 U; 0 Other;
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AAV97774/C
ID AAV97774 standard; RNA; 17 BP.
X
                                                                                                                                                    1 UCGGGUGUCUGCUCUC 17
                                                                                                                                                                                                                                                                           AAT76486 standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                            (first entry)
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ses 14; Conserv
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36-JUN-1996; 17-JUN-1995;

WO9640162-A1

Synthetic.

19-DEC-1996

16-SEP-1997

AAT76486;

RESULT 737 AAT76486/0

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Gaps

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Query Match Best Loca Matches

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The present invention describes enzymatic nucleic acid molecules (NAMs) which specifically cleave RNA derived from an epidermal growth factor receptor (EGF-R) gene. AAV9721 to AAV980403 and AAV98979 to AAV99090 represent specifically claimed target sequence from human EGF-R. AAV98044 to AAV98866 and AAV98867 to V9978 represent hammerhead ribozymes and hairpin ribozymes respectively for human EGF-R. The NAMs are useful for cleaving EGF-R RNA in the treatment of a condition associated with EGFR expression levels e.g. to inhibit cell proliferation in the prevention or treatment of cancers. The NAMs can also be used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of EGF-R RNA in a cell growth factor beta-1; TGF beta-1; antisense oligonucleotide; Enzymatic nucleic acids - which cleave RNA derived from an epidermal growth factor receptor, useful for inhibiting cell proliferation and for / Match 2.9%; Score 12.2; DB 1; Length 17; Local Similarity 82.4%; Pred. No. 4.2e+02; les 14; Conservative 0; Mismatches 3; Indels TGP-beta-1 antisense oligonucleotide TGF-beta1-31. Sequence 17 BP; 3 A; 6 C; 5 G; 0 T; 3 U; 0 Other; (BIOG-) BIOGNOSTIK GES BIOMOLEKULARE DIAGNOSTIK. Claim 5; Page 79; 109pp; English. Akhtar S, Fell P, Mcswiggen JA 391 GCGCCAAGAAGGTCTTC 407 AAV48482 standard; DNA; 17 BP. modulate; gene expression; ss 97EP-00101531. 97EP-00101531. 17 dedechrahadecerre 1 98WO-US000730 97US-0036476P. 97US-00985162. (RIBO-) RIBOZYME PHARM INC. (UYAS-) UNIV ASTON. 15-OCT-1998 (first entry) WPI; 1998-437449/37 treating cancers. Transforming Homo sapiens 31-JAN-1997; 31-JAN-1997; 31-JAN-1997; WO9833893-A2 14-JAN-1998; 04-DEC-1997; EP856579-A1 05-AUG-1998 06-AUG-1998 Synthetic Query Match AAV48482; 740 Matches AAV48482, RESULT XIXBXBXBXBXBXBXBXBXBXBXC ઠે g

WPI; 1998-400910/35.

Preparation of antisense oligo:nucleotide(s) which lack long runs of consecutive guanosine or inosine - and have specific ratio of residues able to form two or three hydrogen bonds, have greater activity and reduced toxicity, used therapeutically or to modulate growth of cells in culture

Claim 10; Fig 3b; 286pp; English

transforming growth factor beta-1 (TGF beta-1). The oligonucleotides exemplify the invention. The specification describes oligonucleotides that contain 8-30 nucleotides, which contain at most 8 nucleotides that contain 8-30 nucleotides, which contain at most 8 nucleotides that consecutive nucleotides able to form three H-bonds each to four consecutive organises, do not contain two sequences of three consecutive consecutive organises, and the ratio between residues able to form two H-bonds each consecutive cytosines, and the ratio between residues able to form two H-bonds each (ZR) or three such bonds (3R) is given by 2R/3R = 0.33-0.72. The oligonucleotides are used to modulate expression of genes, particularly proliferation of primary cell cultures (e.g. bone marrow stem, liver or kidney cells, osteoclasts, osteoblasts and/or keratinocytes). The coligonucleotides can also be used to analyse function of proteins (by cancer or (targeting TGF) for stimulating the immune system cases

Sequence 17 BP; 6 A; 3 C; 5 G; 3 T; 0 U; 0 Other;

Query Match
2.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 82.4%; Pred. No. 4.2e+02;
Matches 14; Conservative 0; Mismatches 3; Indels 0;

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350 GCTCTACAGCGACTTCC 366 **GCTGTACATTGACTTCC** 17 ઠે

RESULT 741 AAX06941

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Gaps ö

AAX06941 standard; DNA; 17

AAX06941;

(first entry) 10-MAY-1999 Canine factor VIII gene fragment PCR primer CS-nt-UTR-U

Factor VIII; canine; dog; diagnosis; animal model; haemophilia A; gene therapy; PCR; primer; 88.

Canis familiaris. Synthetic

CA2225189-A.

06-SEP-1998

98CA-02225189. 06-MAR-1998; 97US-0039953P. 98US-00035141. 06-MAR-1997; 05-MAR-1998;

(TOOH) UNIV QUEENS KINGSTON.

Hough Horrocks L, ບັ Cameron C, Notley Lillicrap D,

WPI; 1999-071205/07.

New canine factor VIII polynucleotide and polypeptide - useful for detection and treatment of haemophilia A using gene therapy.

Schlingensiepen K, Brysch W;

Wed Apr 21 12:58:21 2004

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This is the nucleotide sequence of canine factor VIII gene fragment 6A-3 first round PCR primer CS-nt-UTR-U, where CS indicates canine-specific, and U refers to the amplification region being upstream of the primer. The canine factor VIII gene nucleotide sequence (see AAV99801) was obtained by concatamerisation of RT-PCR-amplified factor VIII fragments obtained from canine liver total RNA (see AAX06886-918), and the sequence was confirmed by RT-PCR (see AAX06919-41). The invention also provides canine factor VIII polypeptides (see AAW80989) and methods for the detection and treatment of canine disorders characterised by factor VIII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 deficiency, especially haemophilia A
Example 2; Page 57; 153pp; English
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Seguence 17 BP; 2 A; 7 C; 5 G; 3 T; 0 U; 0 Other;

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Gaps
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 2.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 4.2e+02; vative 0; Mismatches 3; Indels
Cuery Match
Best Local Similarity 82.4'
Matches 14; Conservative
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RESULT 742 AAV91040

AAV91040 standard; RNA; 17 BP. AAV91040;

Human C-raf target site nucleotide position 747. 18-FEB-1999 (first entry)

Human; c-raf; A-raf; B-raf; hammerhead ribozyme; hairpin ribozyme; target; substrate; catalyst; modulation; expression; Raf gene; delivery; screening; identification; synthesis; deprotection; purification; cancer; inflammation; psoriaals; non-hepatic ascites; infection; genetic drift; restenosis; rheumatoid arthritis; ss.

Homo sapiens

WO9850530-A2

2-NOV-1998

98WO-US009249

35-MAY-1998;

97US-00460S9P. 97US-0049002P. 97US-0051718P. 97US-0056808P. 97US-0061321P. 97US-0061324P. 97US-0068212P. 22-AUG-1997; 02-OCT-1997; 02-OCT-1997; 05-NOV-1997; 39-MAY-1997; 09-NUT-1997

(RIBO-) RIBOZYME PHARM INC.

19-DEC-1997;

Bellon L; Burgin A; Matulic-Adamic J, Reynolds M, Kisich K, Beigelman L, Mcswiggen JA, Karpeisky A, J, Workman CT, Beaudry A, Sweedler D; Thompson J, Jarvis T, Parry T, 1

WPI; 1999-009494/01.

Identifying new catalytic mucleic acid that modulates selected processes - especially ribozymes that cleave Raf RNA for treating cancer, restenosis, and also new ribozymes and modified nucleoside triphosphates used as antiviral agents and synthons.

Claim 177; Page 148; 259pp; English.

A method has been developed for the identification of a nucleic acid

comprises: (a) introducing into the system a random library of nucleic comprises: (a) introducing into the system a random library of nucleic acid catalysts (NAC) having a substrate binding domain (SBD). comprisaing a random sequence, and a catalytic domain (CD); and (b) identifying NAC in systems where modulation has occurred and/or determining the sequence of at least part of the SBDs in such systems. Nucleic acid molecules with cendonuclease activity and catalytic activity, from the present invention, are used to modulate gene expression in plant and mammalian cells and to cleave target nucleic acid, particularly for treating systemic diseases cleaved by specific RNA, e.g. cancer, inflammation, psoriasis, non-hepptic ascites and infection. They may also be used to detect genetic drift and solutions in diseased cells and to determine c-raf RNA. Specifically NACs with RNA-cleaving activity that modulate expression of the Raf gene, are used to treat cancer, restenosis, psoriasis or rheumatoid arthritis, or used to treat cancer, restenosis, psoriasis or rheumatoid arthritis, or of sugar/phosphate modifications in discases stability against nuclease and activity. AAV99922 to AAV93877 represent NACs that can be used in the method, specifically for modulating the expression of a Raf gene

Sequence 17 BP; 4 A; 5 C; 5 G; 0 T; 3 U; 0 Other;

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Gaps
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     2.9%; Score 12.2; DB 1; Length 17; 70.6%; Pred. No. 4.2e+02; Azive 2; Mismatches 3; Indels
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56 AGAGGAGTCTCTGCACT 72 Aguddaduccaacadu 17

ò 요 RESULT 743 AAV92615/

AAV92615 standard; RNA; 17

AAV92615;

(first entry) 18-FEB-1999 Human A-Raf substrate position 2094.

Human; c-raf; A-raf; B-raf; hammerhead ribozyme; hairpin ribozyme; target; substrate; catalyst; modulation; expression; Raf gene; delivery; screening; identification; synthesis; deprotection; purification; cancer; inflammation; psoriasis; non-hepatic ascites; infection; genetic drift; restenosis; rheumatoid arthritis; ss.

Homo sapiens.

WO9850530-A2

12-NOV-1998

98WO-US009249 05-MAY-1998; 09-MAY-1997; 09-JUN-1997

97US-0046059P. 97US-0049002P. 97US-0056808P. 97US-0061321P. 97US-0061324P. 97US-006486EP. 97US-0068212P. 02-OCT-1997 22-AUG-1997 05-NOV-1997

(RIBO-) RIBOZYME PHARM INC

19-DEC-1997;

Bellon L; Burgin A; Jarvis T, Matulic-Adamic J, Reynolds M, Kisich K, Parry T, Beigelman L, Mcswiggen JA, Karpeisky A, Thompson J, Workman CT, Beaudry A, Sweedler D;

WPI; 1999-009494/01.

Identifying new catalytic nucleic acid that modulates selected - especially ribozymes that cleave Raf RNA for treating cancer,

Page 381

restenosis, and also new ribozymes and modified nucleoside triphosphates used as antiviral agents and synthons. Claim 177; Page 161; 259pp; English ##X#X666666666666666666666666

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capable of modulating a process in a biological system. The method capable of modulating a process in a biological system. The method capable of modulating a process in a biological system. The method comprises: (a) introducing into the system a random library of nucleic acid catalysts (NAC) having a substrate binding domain (SBD), comprising a random sequence, and a catalytic domain (CD); and (b) identifying NAC in systems where modulation has occurred and/or determining the sequence of at least part of the SBDs in such systems. Nucleic acid molecules with cadounclease activity and catalytic activity, from the present invention, are used to modulate gene expression in plant and ammalian cells and to cleave target nucleic acid, particularly for treating systemic diseases caused by specific RNA, e.g. cancer, inflammation, psoriasis, non-hepptic ascites and infection. They may also be used to detect genetic drift and matained in the RNA-cleaving activity that modulate expression of the Raf gene, are used to treat cancer, restenosis, psoriasis or rheumatoid arthritis, or generally any condition associated with the level of c-raf. Introduction of sequence activity. AAV90922 to AAV93877 represent NACs that can be used in the activity. AAV90922 to AAV93877 represent NACs that can be used in the continued activity. Accounted the expression of a Raf gene
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Match 2.9%; Score 12.2; DB 1; Length 17; Local Similarity 82.4%; Pred. No. 4.2e+02; les 14; Conservative 0; Mismatches 3; Indels Sequence 17 BP; 1 A; 6 C; 3 G; 0 T; 7 U; 0 Other; 28 AGGGCTGGGACGAAGAT 44 17 AGGCAGAGÁCGAACAT 1 Best Loca Matches

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0; Gaps

Endothelial nitric oxide synthase antisense oligonucleotide. AAX54277 standard; DNA; 17 BP. (first entry) 05-JUL-1999 AAX54277; RESULT 744 AAX54277,

Antisense oligonucleotide; multiple target; antisense treatment; impaired respiration; inflammation; lung disease; pulmorary vasconstriction; inflammation; allergic rhinitis; acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; pulmonary disease; leukemia; imphysema; chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pancreatic cancer; hepatocellular carcinoma; kidney cancer; melanoma; hepatic metastasis; prostate cancer; ss

409913886-A1 Synthetic.

25-MAR-1999.

98WO-US019419. 17-SEP-1998;

97US-0059160P. 17-SEP-1997; 09-JUN-1998;

(UYEC-) UNIV EAST CAROLINA.

vyce JW;

WPI; 1999-229400/19

New antisense oligonucleotides used in treatment of, e.g. pulmonary vasoconstriction.

Disclosure, Page 61; 120pp; English.

The specification describes antisense oligonucleotides (AAX52869-X55271)

directed against at least 2 mRNAs selected from target genes, coding and

con-coding regions of RNAs corresponding to target genes, coding and

codons, genomic flanking regions, intron-exon borders, the 5' end, the 3'

codons, genomic flanking regions, intron-exon borders, the 5' end, the 3'

coditions or mixtures. The antisense oligonucleotides may be derived

cronditions or mixtures. The antisense oligonucleotides may be derived

from sequences AAX55272-74. These multiple target oligonucleotides

conditions or mixtures. The multiple target oligonucleotides

conditions or mixtures. The multiple target oligonucleotides

conditions or mixtures. These multiple target oligonucleotides

conditions or mixtures. Typical diseases and conditions are those associated with impaired respiration and inflammation, including lung diseases, pulmonary vasoconstriction, inflammation, allergic rhinitis, catter asthma, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, confirmation, engiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukemias, lymphomas, carcinomas e.g.

colon cancer, breast cancer, lung cancer, pancreatic cancer, melanoma, hepatic metastassized well as all types of cancers which may metastasize or have metastasized to the lungs, including breast and prostate cancer

Sequence 17 BP; 0 A; 7 C; 6 G; 4 T; 0 U; 0 Other;

Query Match
2.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 82.4%; Pred. No. 4.2e+02;
Matches 14; Conservative 0; Mismatches 3; Indels

268 ACCTGGAGCAGGCGGC 284

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AAX29695 standard; DNA; 17 BP. AAX29695; RESULT 745 AAX29695,

Human bone morphogenic protein (BMP)-2 forward primer. 04-JUN-1999 (first entry)

BMP; BMP-2; bone morphogenetic protein; tissue regeneration; skin; bone; cartilage; tendon; ligament; muscle; connective tissue; nerve; cardiac; liver; lung; kidney; pancreas; brain; embryonic development; growth factor; osteoporosis; osteoarthritis; fracture; PCR primer; ss.

Homo sapiens.

98WO-US018603. WO9911664-A1. 04-SEP-1998; 11-MAR-1999. THE KEY TO THE WAR TO THE TOTAL TO THE TOTAL TO THE TANK TO THE TANK TO THE THE TA

97US-0057989P. 98US-00148234. (GEMY) GENETICS INST INC. (YISS) YISSUM RES & DEV CO. 05-SEP-1997;

Turgeman G; Moutsatsos I, Gazit D, Zilberman Y,

WPI; 1999-214697/18.

Production of cells for implantation at the site of bone infirmity in a human, using DNA encoding a bone morphogenetic protein - useful for

treating osteoporosis, osteoarthritis and non-union fractures.

Example 14; Page 43; 71pp; English

The invention relates to the production of oils for implantation at the site of a bone infirmity in a human, that comprises transforming and culturing a host containing DNA encoding a bone morphogenetic protein (EMP). The method is useful for regenerating various tissues, including bone, cartilage, tendon, ligament, muscle, skin (and other connective tissues), nerve, cardiac, liver, lung, kidney, pancreas, and brain. The method is also useful for inducing and/or regeneration of tissue, including the induction of epidermal, endodermal and mesodermal tissue including the induction of epidermal, endodermal and mesodermal tissue including the induction of epidermal, endodermal and mesodermal tissue including the induction of epidermal, endodermal and mesodermal tissue including the number of growth factors broduced by the method are useful for treating osteoporasis and osteoarthritis, non-union can be used for growth factor delivery to signalling receptors of transplanted cells (autocrine effect) and host progenitor stem cells (paracorine effect) for the engraftment, differentiation, and stimulation of new bone growth. Therefore, the method provides an effective therapy for non-union fractures. Sequences AAX29695-696 represent primers for BMP

Sequence 17 BP; 3 A; 9 C; 2 G; 3 T; 0 U; 0 Other;

Ouery Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels 23 GACCGAGGGCTGGGACG 39

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Gaps

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17 Grcadaddcrdddard 1 g

AAA33721 standard; DNA; 17 BP AAA33721/0

(first entry) 28-JUL-2000 Low adenosine antisense oligonucleotide SEQ ID NO:1410.

Human, adenosine receptor; low adenosine antisense oligonucleotide; phosphorothioate; impaired respiration; inflammation; allergy; allergy; disease; bronchoconstriction; inhibitor; antihinflammatory; antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway; lung disease; ischaemic condition; pulmonary vasoconstriction; asthma; respiratory distress syndrome; pain, cystic fibrosis; emphysema; pulmonary hypertension; chronic obstructive pulmonary disease; COPD; cancer; leukaemia; lymphoma; carcinoma; metastasis; se.

lomo sapiens.

VO200009525-A2.

24-FEB-2000.

3-AUG-1999;

98US-0095212P. 3-AUG-1998;

99WO-US017712

UYEC-) UNIV EAST CAROLINA.

Nyce JW;

WPI; 2000-205971/18.

New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstruction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or

Claim 18; Page 441; 1343pp; English.

The present invention describes a new composition couplisation conjugation describes a new composition call introduced in bronchoconstriction, allergies, and/or inflammation. The Ow can have antiinflammatory, antiallergies and/or inflammation. The Own can have antiinflammatory, antiallergies and/or contasthmatic, cytosteatic and analgesic activities. The compositions are useful for the treatment of diseases associated with inflammation, impaired airways, including lung diseases and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating e.g. ischaemic conditions, pulmonary vasoconstriction, allergies, asthma, important conditions, pulmonary vasoconstriction, allergies, asthma, inparable in conditions, pulmonary vasoconstriction, allergies, asthma, pulmonary disease (COPD), and cancers such as leukaemias, lymphomas, or arcinomas, and cancers which may metastasise to the lungs, including breast and prostate cancer. The A-containing ONS break down with the own of deoxyadenosine which activates adenosine receptors causing bronchoconstriction and inflammation. AAA32313 to AAA35312 represent to nucleotide sequences given in the sequence listing from the present correspond to SEQ ID NO:11 to 1885, but the sequences differ from the previously named sequences. SEQ ID NO:11 to 1680 (AAA3233 to AAA3332) to AAA33313 to AAA33313 to AAA33313 to AAA33313 to AAA33313 to AAA33333 to AAA3333 to AAA33333 to AAAA33333 to AAA33333 to AAA33333 to AAA33333 to AAA33333 to AAA333

Sequence 17 BP; 0 A; 7 C; 6 G; 4 T; 0 U; 0 Other;

3; Indels 0; Query Match
2.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 82.4%; Pred. No. 4.2e+02;
Matches 14; Conservative 0; Mismatches 3; Indel8

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268 ACCTGGAGCAGGCGGC 284

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AAA91985 standard; DNA; 17 BP. RESULT 747 AAA91985

(first entry) 10-JAN-2001

AAA91985;

Nested PCR primer 1184F for S. neurona small ribosomal subunit

Small ribosomal subunit; SRSU; Equine protozoal myeloencephalitis; EPM; diagnosis; nested PCR primer; ss.

Sarcocystis neurona.

29-AUG-2000.

95US-00388029. 14-FEB-1995; 14-FEB-1995;

(KENT) UNIV KENTUCKY RES FOUND.

Fenger CK, Gajadhar AA, Dubey JP,

Granstrom DE;

WPI; 2000-586347/55.

Sarcocystis neurona diagnostic primer, useful for in vitro diagnostic testing for Equine protozoal myeloencephalitis, i.e. for diagnosing the presence of S. neurona in equine blood or cerebrospinal fluid.

Example 3; Col 7; 41pp; English

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The present invention relates to a diagnostic primer from positions 1470-1487 of the small ribosomal subunit of Sarcocystis neurona. This primer is unique to the S. neurona species. The primer is useful for diagnostic tests for Equine protozoal myeloencephalitis (EPM) where the presence of S. neurona is indicative of EPM. The present sequence is a nested PCR primer used in the diagnostic assay to identify S. neurona
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Sequence 17 BP; 2 A; 5 C; 8 G; 2 T; 0 U; 0 Other;

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Gaps
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0
Score 12.2; DB 1; Length 17; Pred. No. 4.2e+02; 0; Mismatches 3; Indels
 Query Match
2.9%; Soc
Best Local Similarity 82.4%; Pro
Matches 14; Conservative 0;
                                                                  3 CCAGGAGTGAACTGCG 19
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1 CCAGGCGTGGAGCTGCG 17

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AAZS6635 standard; DNA; 17 BP. AAZ56635; RESULT 74

21-MAR-2000 (first entry)

Canine Factor VIII isolation and cloning PCR primer SEQ ID NO:61. Canine; factor VIII; haemostatic; diagnosis; haemophilia A; dog; PCR primer; ss.

Canis sp.

CA2264431-A1

05-SEP-1999

99CA-02264431. 05-MAR-1999;

98US-00035141. 98CA-02225189. 05-MAR-1998; 06-MAR-1998; (TOOH) UNIV QUEENS KINGSTON

Cameron C; Lillicrap D, C, Notley C, Hough WPI; 2000-073270/07. Horrocks LSH,

Isolated nucleic acid encoding a canine factor VIII polypeptide for treating a disorder characterized by canine factor VIII deficiency, such as hemophilia A.

Example 2; Page 58; 152pp; English.

The present invention describes canine factor VIII. The isolated factor VIII nucleic acid molecule and protein can be used for treating a disorder characterised by canine factor VIII defliciency in a canine, especially haemophilia A. AAZ56579 to AAZ5603s represent primers used in the isolation and cloning of canine factor VIII

Sequence 17 BP; 2 A; 7 C; 5 G; 3 T; 0 U; 0 Other;

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Gaps
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Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels
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214 AGAACTCGGTGGCGGCC 230 17 AGACCTCGCTGTCGGCC

ઠે 임 RESULT 749

ВЪ.

Human endothelial nitric oxide synthase polynucleotide fragment #1410

human; airway disorder; bronchoconstriction; lung inflammation; human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory, bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; crespiratory obstruction; pulmonary obstruction; immeded respiration; respiratory obstruction; pulmonary vasconstriction; asthma; RDS; respiratory distress syndrome; pain; cystic fibrosis; allergic rhinitis; pulmonary hypertension; emphysema; pain, cystic fibrosis; allergic rhinitis; chronic obstructive pulmonary disease; pulmonary infection; bronchitis;

24-MAR-2000; 2000WO-US008020.

99US-0127958P.

Low adenosine (A) content antisense oligonucleotides which do not trigg adenosine receptors during metabolism, useful e.g. for treating cancers and respiratory obstructions.

Claim 14; Page 251; 1592pp; English.

The present invention describes low adenosine (A) content antisense oligonucleotides and compositions (I) comprising them. In the antisense oligonucleotides the A is replaced by a 'Universal' or alternative base. (I) can have respiratory, bronchodilator, antiinflammatory, analgesic, immunosuppressive, antiatehmatic, hypotensive and cytostatic activities. The antisense oligonucleotides and (I) can be used to down-regulate the expression and or activity of target polypeptides associated with communication proteins and malignancies, such as stimulating and activating peptide factors and transmitters, transcription factors, immunoglobulins and antibodies, antibody receptors, cytokines and chemokines, endogenously produced specific and non-specific enzymes, chemokines, endogenously produced specific and non-specific enzymes, binding proteins, adheosine receptors, bradykinin receptors, cremokine receptors, adenosine receptors, bradykinin receptors cytokines and chemokine receptors, binding proteins and malignancy associated proteins. The receptors, binding proteins and malignancy associated proteins. The antisense oligonucleotides may be used in this way to treat disorders including respiratory obstruction (especially bulmonary obstruction (especially bulmonary obstruction and/or bronchoconstriction) and/or lung inflammation, allergic and/or bronchoconstriction, and associated with a disease or condition selected from pulmonary vasoconstriction, inflammation, allergic explanner, condition specially espiratory distress syndrome (RDS), pain, cystic fibrosis (CF), allergic rhinitis (AR), pulmonary conditions plected from pulmonary vasoconstriction, inflammation, and/or horselfice and/or exemplification pelection, bulmonary infections, bronchitis, and/or the present invention

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Wed Apr 21 12:58:21 2004
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Matches

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The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor. EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and
                                                                                                                                                                                                                                                                                                                                                                                                    Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ribozyme; erythropoietin; granulocyte colony stimulating factor;
interferon alpha; ss.
                                                    Ribozyme; erythropoietin; granulocyte colony stimulating factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 4.2e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 0 A; 7 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                 Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hammerhead ribozyme substrate #3502.
                   Hammerhead ribozyme substrate #879.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 37; Page 76; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               240 GGCTGCTTCCCGGGCTC 256
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                                                                                                                                                                                                                                                                                                                               Blatt L, Zwick M, Pavco P,
                                                                                                                                                                                                                     11-APR-2000; 2000WO-US009721
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                                                                                                                                                                                                                                                                                            (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAF07245 standard; DNA; 17
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les 14; Conservative
                                                                      interferon alpha; ss.
                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-647423/62
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                                                                                                                                             WO200061729-A2
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                                                                                                             Homo sapiens
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
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                                   Length 17;
                                 ch 2.9%; Score 12.2; DB 1; Length 1
1 Similarity 82.4%; Pred. No. 4.2e+02;
14; Conservative 0; Mismatches 3; Indels
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Sequence 17 BP; 0 A; 7 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Blatt L, Zwick M, Pavco P, McBwiggen J;
                                                                                                                                                                                                                                                                                                                                                 Hammerhead ribozyme substrate #2500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 18; Page 113; 164pp; English.
                                                                                                         268 ACCTGGAGCAGGCGGC 284
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AAF05281 standard; DNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                      interferon alpha; ss.
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Best Loca Matches

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RESULT 751

AAF02584
ID AAF0
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AC AAF0
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DT 16-F

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Gaps

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Sequence 17 BP; 2 A; 6 C; 4 G; 5 T; 0 U; 0 Other; Blatt L,

The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoletin, granulocyte colony stimulating factor protein and interferon alpha Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin. Zwick M, Pavco P, Mcswiggen J; Claim 54; Page 136; 164pp; English. WPI; 2000-647423/62

Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indel8 167 GGTGTACTACGAGTCCA 183 GGTGTTCTACCCGTCCA 17 ò g

Gaps

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AAF05334 standard; DNA; 17 (first entry) 16-FEB-2001 AAF05334; RESULT 753 AAF05334

BP.

Hammerhead ribozyme substrate #2553.

Ribozyme; erythropoietin; granulocyte colony stimulating factor; 11-APR-2000; 2000WO-US009721. interferon alpha; ss. WO200061729-A2. Homo sapiens. 19-OCT-2000

99US-0129390P (RIBO-) RIBOZYME PHARM INC. 12-APR-1999;

Blatt L, Zwick M, Pavco P, Mcswiggen J;

WPI; 2000-647423/62.

Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.

Claim 18; Page 114; 164pp; English.

The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoletin, granulocyte colony stimulating factor protein and interferon alpha

Gaps .. 2.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 4.2e+02; tive 0; Mismatches 3; Indels Sequence 17 BP; 1 A; 7 C; 4 G; 5 T; 0 U; 0 Other; Best Local Similarity 82.4 Matches 14; Conservative Query Match

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ABK01641 standard; RNA; 17 RESULT 754 ABK01641/c ID ABK0164

12-MAR-2002 (first entry) ABK01641;

Human NOGO G-Cleaver #97.

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; noctropic; neuroprotective; antiparkinsonian; muscular; D20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; g-cleaver; amberzyme; inizyme; lumphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphomy; leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MC; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; cerebrovascular accident; gamotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntingcon's disease; central capabenetative disease. Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.

sapiens Synthetic. Homo

WO200159103-A2.

16-AUG-2001.

09-FEB-2001; 2001WO-US004273

11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P.

(RIBO-) RIBOZYME PHARM INC. (BLAT/) BLATT L. (MCSW/) MCSHIGGEN J. (CHOW/) CHOWRIRA B M.

Chowrira BM; Blatt L, Mcswiggen J,

WPI; 2001-607195/69.

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Claim 88; Page 93; 200pp; English.

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NOGO). The nucleic acids (e.g. arbbzyme or a DNAzyme) an Inozyme (an endolytic nucleic acids (e.g. arbbzyme or a DNAzyme) an INOZYME (aleaving RNA with a NYN motif) pran ambersyme (cleaving RNA with a NYN motif) pran an ambersyme (cleaving RNA with a NYN motif) pran an arbarsyme (cleaving RNA with an NGN triplet), a zinzyme (cleaving RNA with a YGY motif). The CD20-tergetting nucleic acid is used to cleave RNA of CD20 in the presence of a divalent cation that is preferably Mg^2^+.

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CC Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level CC of CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-cc treat lymphoma (MLL), bulky low-grade or follicular MLL, lymphocytic lymphoma (MCL), immunocytona (MC), small B-cell lymphocytic lymphoma, cc lymphoma (MCL), immunocytona (MC), small B-cell lymphocytic lymphoma, immunocytopaenia, and inflammatory arthropathy. The NOGO-targetting nucleic acid is used to cleave RNA of the NOGO gene in the presence of a divalent cation that is preferably MG'2+. Furthermore, the nucleic acid may be contacted with a cell to reduce NOGO activity of the conspice a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more therapies. In particular, the NOGO-targetting nucleic acid may be used to thearapies. In particular, the NOGO-targetting nucleic acid may be used to contential nervous system (CNS) injury and cerebrosscular accident (CNA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS), chemotherapy-induced neuropathy, and/co there neurodegenerative disease stackia, muscular dystrophy, and/co to ther neurodegenerative disease stackia, muscular dystrophy, and/co to ther neurodegenerative disease stackia, muscular of the modulation of NOGO expression. The present content and acceled the modulation of the invention

Sequence 17 BP; 5 A; 2 C; 7 G; 0 T; 3 U; 0 Other;

Gaps ö 2.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 4.2e+02; ative 0; Mismatches 3; Indels Query Match
Best Local Similarity 82.4'
Matches 14; Conservative

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RESULT 75 ABK02370

ABK02370 standard; RNA; 17 BP. (first entry) 12-MAR-2002 ABK02370;

Human NOGO Amberzyme #42.

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurice growth inhibitor gene; NOGO; hammerhead ribozyme; bnAzyme; inozyme; dela growth inhibitor gene; NOGO; hammerhead ribozyme; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; MC; immunocytoma; non-Hodgkin's lymphoma; NHC; immunocytoma; stroke; dementia; inflammatory arthropathy; central nervous system injury; dementia; inflammatory arthropathy; central nervous system injury; cheentherapy-induced neuropathy; amyotrophic lateral sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; parkinson; disease; multiple sclerosis; parkinson; disease; ataxia; Huntington's disease; Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.

sapiens. Synthetic. JOHO

40200159103-A2.

16-AUG-2001.

39-FEB-2001; 2001WO-US004273.

11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P.

RIBOZYME PHARM INC. BLATT L. MCSWIGGEN J. (RIBO-) H (BLAT/) H (MCSW/) N

(CHOW/) CHOWRIRA B M.

Chowrira BM; Mcswiggen J, Blatt L,

WPI; 2001-607195/69.

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Claim 88; Page 131; 200pp; English.

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down exprassion of a neurite growth inhibitor gene (NGO). The regulates expression of a neurite growth inhibitor gene (NGO). The nucleic acids (e.g. a ribozyme or a numberzyme (cleaving RNA with a NNY motif) properties of CD20 the claim of the presence of a divalent cation that is preferably MG²+. Of CD20 in the presence of a divalent cation that is preferably MG²+. Of CD20 in the presence of a divalent cation that is preferably MG²+. Of CD20. The treatment may further comprise the use of one or more che cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more checken is In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular nontreledgating nucleic acid may be contacted with a cell to reduce NOSO gene in the craspetting nucleic acid may be contacted with a cell to reduce NOSO gene in the cell and treat a patient having a condition associated with the level of the condition associated with the level of the contacted with a cell to reduce NOSO gene in the cell and treat a patient having a condition associated with the level of the condition associated with the level of the cell and treat a patient having a condition associated with the level of the condition associated with the level of the cell and treat a patient having a condition associated with the level of the cell and treat a patient having a condition associated with the level of therapies. In particular, which will be seen the cell of the cell sequence is an amberzyme molecule of the invention

Sequence 17 BP; 7 A; 2 C; 8 G; 0 T; 0 U; 0 Other;

ö Query Match
2.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 82.4%; Pred. No. 4.2e+02;
Matches 14; Conservative 0; Mismatches 3; Indels

27 GAGGGCTGGGACGAAGA 43 1 caccaccaccaccaccaca 17 ò a

ABA81116 standard; DNA; 17 BP. ABA81116

RESULT 756

ABA81116;

24-JAN-2002 (first entry)

UGT1 mutation correcting oligonucleotide SEQ ID NO: 3962.

Human; gene therapy, adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MIH1; APOE; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hypercholesterolaemia; UGH1; syndrome; APP; PSEN1; antisense;

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The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCAL, BRCAZ, CFTR, cyclin-dependent kinase inhibitor 2A (CDKN2A), APC, Factor VIII, Factor IX, haemoglobin alpha locus (CDKN2A), hemoglobin alpha locus 2 (HBA2), MLHI, MSH2, MSH6, apolipoprotein B (APDE), LDL receptor (LDLR), UDP-glucuronosyltransferase (UGTI), amyloid precursor protein (APC), presenilin-2 (PSEN2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, various syndromes. The present sequence is one of the colon and various syndromes. The present sequence is one of the gene correcting oligonucleotides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2, CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent Kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MIH1; APOB; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein E; LDLR;
UDP-glucuronosyltransferase, amyloid precursor protein; presenilin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic; antilipemic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 7; Page 258; 294pp; English.
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27-MAR-2000; 2000US-019219P.
UJUN-2000; 2000US-0206538P.
30-OCT-2000; 2000US-024989P.
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modification
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The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the coligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, cretinoblastome, BRCAI, BRCAZ, CFTR, CYCLIn-dependent kinase inhibitor 2A cobnigonary APC, BCCtor V, Factor VIII, Factor IX, haemoglobin alpha locus (HBAI), haemoglobin alpha locus 1 (HBAI), haemoglobin alpha locus 1 (HBAI), haemoglobin alpha locus 1 (HBAI), amyloid precursor protein (APC), presentlin-1 (BSENI) and precursor protein (APC), presentlin-1 (PSENI) and precursor protein (APC), presentlin-1 (PSENI) and precursor meaning deficiency, cystic fibrosis, such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemoghila, hypercholesterolaemia, thalassaemia, sickle cell anaemia, all all anaemia, all anaem
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familial hypercholesterolaemia, UGT1; syndrome; APP; PSEN1; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presenilin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and for chemical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide for targeted alterations of genetic sequences treating cystic fibrosis, comprises at least one mismatch and modification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 4.2e+02; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LDLR mutation correcting oligonucleotide SEQ ID NO: 3695.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 3 A; 4 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     various syndromes. The present secoligonucleotides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 7; Page 44; 294pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rice MC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      멾.
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27-MAR-2000; 2000US-0192179F.
10-UTN3-2000; 2000US-0208538F.
30-OCT-2000; 2000US-0244989F.
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Best Local Similarity 82.2.2
-1.58 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gamper HB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UYDE ) UNIV DELAWARE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-639230/73.
                                                                                  antilipemic; ss
                                                                                                                                                                                        WO200173002-A2
                                                                                                                                        Homo sapiens
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The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, criticoblastoma, BRCAL, BRCAL, CFTR, CYCLIn-dependent Kinase inhibitor 2A retinoblastoma, BRCAL, BRCAL, CFTR, CYCLIn-dependent Kinase inhibitor 2A (CDXRZA), APC, Factor V, Pactor VIII, Factor IX, haemoglobin alpha locus in (HBA1), haemoglobin alpha locus 2 (HBA2), MLH1, MSH2, MSH6, apolipoprotein E (APDC), LD. receptor (LDLR), UD-glucuronosyltransferase (UGT1), amyloid precursor protein (APC), presentlin-1 (PSEN1) and presentlin-2 (PSEN2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, chaemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, alzabalmer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting familial hypercholesterolaemia; UGT1; syndrome; APP; PSEN1; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presenilin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic; Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical modification. mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hypercholesterolaemia; UGT1; syndrome; APP; PSEN1; antisense Claim 7; Page 245; 294pp; English. oligonuclectides of the invention Rice MC 27-MAR-2000; 2000US-0192176P. 27-MAR-2000; 2000US-0192179P. UTUN-2000; 2000US-0208538P. 30-OCT-2000; 2000US-0244989P. 27-MAR-2001; 2001WO-US009761. (UYDE) UNIV DELAWARE. Gamper HB, WPI; 2001-639230/73. antilipemic; ss. WO200173002-A2. Homo sapiens. 04-OCT-2001. Kmiec EB,

; 0 Gaps ö 2.9%; Score 12.2; DB 1; Length 17; llarity 82.4%; Pred. No. 4.2e+02; Conservative 0; Mismatches 3; Indels Sequence 17 BP; 4 A; 4 C; 9 G; 0 T; 0 U; 0 Other; Local Similarity tes 14; Conserv Query Match Best Loc Matches

90 CGAAGGCCGAGCAGGGG 17 CGAGGCCGCGCAGTGG 74 ð

ABA81117 standard; DNA; 17 BP ABA81117; RESULT 759 **ABA**8111 **EZZZZZZZZZZ**

(first entry) 24-JAN-2002

UGT1 mutation correcting oligonucleotide SEQ ID NO: 3963.

Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BFTR; cystic fibrosis; cancer; Bactor V; cyclin-dependent kinase inhibitor 24; CDKN2A; melanoma; APC; HBA1; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis;

The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, can be following genes: adenosine deaminase, p53, beta-globin, hibstor 2A ctrinoblastoma, BRCA1, BRCA2, CFTR, cyclin-dependent kinase inhibitor 2A (CDNXA), APC, Factor VIII, Pactor IX, haemoglobin alpha locus (CDDIA), haemoglobin alpha locus 2 (HBA2), Will, MSH6, MSH6, apolipoprotein E (APOE), LDL receptor (LDLR), UDP-glucuronosyltransferase (UGTI), amyloid precursor protein (APOE), presentlin-1 (PSENI) and presentlin-2 (PSENI). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, hammophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting oligonucleotides of the invention haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOB; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hypercholesterolaemia; UGT1; syndrome; APP; PSEN1; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presentlin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic; antilipemic; ss. Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical modification. Claim 7; Page 258; 294pp; English. Rice MC; 27-MAR-2000; 2000US-0192176P. 27-MAR-2000; 2000US-0192179P. 01-JUN-2000; 2000US-0208538P. 30-OCT-2000; 2000US-0244989P. 27-MAR-2001; 2001WO-US009761 Gamper HB, (UYDE) UNIV DELAWARE WPI; 2001-639230/73. WO200173002-A2. Homo sapiens 04-OCT-2001. Kmiec EB,

ö 2.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 4.2e+02; tive 0; Mismatches 3; Indels 338 CCAGGGCCGGCTGTT 354 14; Conservative Similarity Local Matches ò

Sequence 17 BP; 5 A; 6 C; 6 G; 0 T; 0 U; 0 Other;

Query Match

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Gaps

17

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ABA80848 standard; DNA; 17 (first entry) 24-JAN-2002 ABA80848; ABA80848/C
ID ABA80
XX AC ABA80
XX 24-J.
XY 24-J.
XX BLDLR
XX Huma
XW Huma
XW cycl

RESULT 760

ВЪ.

LDLR mutation correcting oligonucleotide SEQ ID NO: 3694.

Human, gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDRN2A; melanoma; APC; HBA1; HBA2;

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adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein B; LDLR; familial hypercholesterolaemia; UGT1; syndrome; APP; PSSN1; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presentin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic; antilipemic; ss.
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Homo sapiens.

WO200173002-A2.

04-OCT-2001

27-MAR-2001; 2001WO-US009761.

27-MAR-2000; 2000US-0192176P. 27-MAR-2000; 2000US-0192179P. 01-UUN-2000; 2000US-0208538P. 30-OCT-2000; 2000US-0244989P.

(UYDE) UNIV DELAWARE

Kmiec EB, Gamper HB, Rice MC

WPI; 2001-639230/73.

Oligonuclectide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical modification.

Claim 7; Page 245; 294pp; English.

The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CFTR, CYClin-dependent kinase inhibitor 2A (CDKN2A), APC, Factor V, Factor VIII, Factor IX, haemoglobin alpha locus (HBA2), MLH1, MSH2, MSH6, apolipoprotein E (APOES, DL receptor (LDLR), UDP-glucuronosyltransferase (UGT1), amyloid precursor protein (APC), presentlin-2 (PSEN1) and presentlin-2 (PSEN2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, chaemophila, hypercholesterolaemia, thalassaemia, sickle cell anaemia, Alzhaimer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting various syndromes. The present sec oligonuclectides of the invention

Sequence 17 BP; 0 A; 9 C; 4 G; 4 T; 0 U; 0 Other;

.. . Match 2.9%; Score 12.2; DB 1; Length 17; Local Similarity 82.4%; Pred. No. 4.2e+02; les 14; Conservative 0; Mismatches 3; Indels Query Match Best Loca Matches

74 CGAGGGCGGCGCAGTGG 90

17 cgaaggccgagcaggg 1

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RESULT 76 ABA77218

ABA77218 standard; DNA; 17 BP.

ABA77218;

(first entry) 24-JAN-2002

Adenosine deaminase deficiency correcting oligo SEQ ID NO: 64.

Human, gene therapy, adenosine deaminase deficiency, p53; beta-globin, retinoblastoma, BRCA1; BRCA2; CFTR; cystic fibrosis; cancer, Factor V;

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cyciin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBAI; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein B; LDLR; familial hypercholesterolaemia; UGT1; syndrome; APP; PSENI; antisense; UDP-glucuromosyltransferase; amyloid precursor protein; presenilin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic; antilipemic; ss.
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WO200173002-A2.

04-OCT-2001

27-MAR-2001; 2001WO-US009761

27-MAR-2000; 2000US-0192176P. 27-MAR-2000; 2000US-0192179P. 01-UTN-2000; 2000US-0208538P. 30-OCT-2000; 2000US-0244989P.

(UYDE) UNIV DELAWARE,

Rice MC Gamper HB, Kmiec EB,

WPI; 2001-639230/73.

Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical modification.

Claim 7; Page 44; 294pp; English.

The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of ganomic sequences, where the oligonucleotide has at least one mismatch compared with the ganomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaninase, p53, beta-globin, controlled the following genes: adenosine deaninase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CFTR, cyclin-dependent kinase inhibitor 2A controlled to the following genes: adenosine deaninase, p53, beta-globin alpha locus (CDENZA), APC; Ractor VIII, Factor IX, heemoglobin alpha locus (IMBA1), haemoglobin alpha locus 2 (HBA2), mill, MSH2, MSH6, apolipoprotein E (APOE), LDL receptor (LDLR), UDP-glucuronosyltransferase (UGTH), amyloid precursor protein (APC), presentilin-1 (PSENI) and presentilin-2 (PSENS). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, although and although and anticolled as an anticolled and an anticolled controlled controlled colleges of the invention

Sequence 17 BP; 2 A; 8 C; 4 G; 3 T; 0 U; 0 Other;

Gaps ô 2.9%; Score 12.2; DB 1; Length 17; 12.4%; Pred. No. 4.2e+02; ve 0; Mismatches 3; Indels 82.4%; Query Match
Best Local Similarity 82.4

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89 GGACATCACCACGTCTG 105

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ABL47246 standard; RNA; 17 RESULT 762

(first entry) 27-JUN-2003

ABL47246;

Human GRID Amberzyme substrate oligonucleotide #146.

Human; Grb2-related with Insert Domain; GRID; T-cell;

Shannon ME;

Chen W,

Rank DR,

Hanzel DK,

Gu Y, Ji Y, Penn SG, WPI; 2002-179446/23.

(AEOM-) AEOMICA INC.

30-JAN-2001; 2001WO-US000663. 30-JAN-2001; 2001WO-US000664. 30-JAN-2001; 2001WO-US000665. 30-JAN-2001; 2001WO-US000665. 30-JAN-2001; 2001WO-US000668. 30-JAN-2001; 2001WO-US000668. 30-JAN-2001; 2001WO-US000669. 30-JAN-2001; 2001WO-US000669.

The present invention relates to oligonucleotides that downregulate the expression of human Grb2-related with Insert Domain (GRID) gene. GRID is a T-cell co-stimulatory adaptor protein. The oligonucleotides are useful for modulating the expression of GRID, to treat conditions such as tissue/graft rejection and leukaemia. The oligonucleotides can also be administered in conjunction with other therapies such as radiation, Human, genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss. Domain Gaps co-stimulatory adaptor protein; tissue rejection; graft rejection; leukaemia; cytostatic; ss. Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:1480. Ellis JH New nucleic acid(s) for regulating the Grb2-related with Insert D (GRID) gene comprises using antisense and enzymatic nucleic acid molecules such as hammerhead ribozymes. . 0 Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 64.7%; Pred. No. 4.2e+02; Matches 11; Conservative 3; Mismatches 3; Indels Hamblin PA, Sequence 17 BP; 2 A; 2 C; 9 G; 0 T; 4 U; 0 Other; Mcswiggen JA, Claim 4; Page 88; 108pp; English. used to illustrate the invention 289 AGCTGGTGAAGGACCTG 305 1 AGGUGGUGGAGGUCCUG 17 ABN01488 standard; DNA; 17 BP 23-FEB-2001; 2001WO-US005957. 24-FEB-2000; 2000US-0184594P. 21-SEP-2000; 2000US-0234687P. 27-SEP-2000; 2000US-023639P. 04-OCT-2000; 2000GB-00024263. 30-JAN-2001; 2001WO-US000661. 25-MAY-2001; 2001WO-US016981. Jarvis T, Von Carlowitz I, (RIBO-) RIBOZYME PHARM INC. (GLAX) GLAXO GROUP LTD. (first entry) WPI; 2001-550088/61. WO200162911-A2. WO200192524-A2 domo sapiens 30-AUG-2001 ABN01488; RESULT 763 ABN01488/c a ò

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used an gene therapy and vaccine production. The hGDMLP-1 nucleic acids in samples, as amplification substrates and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 and/or amount specifically of hGDMLP-proteins, as specific biomolecule and/or amount supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the ChGDMLP-1 sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO can fire thy wipo.int/pub/published_pot_sequence ö New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1. Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss. 0; Gaps Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:1481. 2.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 4.2e+02; tive 0; Mismatches 3; Indels Sequence 17 BP; 3 A; 5 C; 8 G; 1 T; 0 U; 0 Other; Disclosure, SEQ ID NO 1480, 214pp, English. 125 ceecarecreeceec 141 ABN01489 standard; DNA; 17 BP. 17 ceecricerescasic 1 (first entry) Query Match Best Local Similarity 82.4 Matches 14; Conservative 29-MAY-2002 ABN01489; RESULT 764 ABN01489/c ઠે g

Homo

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 clan be used as probes to detect, characterise and quantify nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specific biomolecule and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as as characterise of for replacement therapy. The production, and in vaccines or for replacement therapy. The production and in vaccines or for replacement therapy. The production and in vaccines or for replacement therapy. The production associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 3 A; 5 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ftp.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 1481; 214pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gu Y, Ji Y, Penn SG, Hanzel DK,
                                                                                                                  26-MAY-2000; 2000US-0207456P.
27-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0234687P.
04-OCT-2000; 2000US-0236359P.
04-OCT-2000; 2001WO-US000661.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
                                                                                                                                                                                                                                                                                                                                                                                           30-JAN-2001; 2001WO-US000670
05-FEB-2001; 2001US-0266860P
                                                                               25-MAY-2001; 2001WO-US016981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-179446/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           (AEOM-) AEOMICA INC.
  WO200192524-A2
                                         06-DEC-2001
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Gaps ö 2.9%; Score 12.2; DB 1; Length 17; 32.4%; Pred. No. 4.2e+02; ved. 0; Mismatches 3; Indels Local Similarity 82.4%; les 14; Conservative Query Match Best Loca Matches

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RESULT 765 ABN06221/C ID ABN06221 standard; DNA; 17 BP. XX AC ABN06221;

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss. Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:6213 25-MAY-2001; 2001WO-US016981 WO200192524-A2. Homo sapiens

(first entry)

29-MAY-2002

27-SEP-2000; 2000US-02363599-04-CCT-2000; 2000US-02365599-04-CCT-2000; 2001WO-US000661. 30-JAN-2001; 2001WO-US000663. 30-JAN-2001; 2001WO-US000663. 30-JAN-2001; 2001WO-US000664. 30-JAN-2001; 2001WO-US000670. 05-FEB-2001; 2001US-0266860P.

Chen W, Shannon ME;

Rank DR,

Chen W, Shannon ME; Gu Y, Ji Y, Penn SG, Hanzel DK, Rank DR, WPI; 2002-179446/23.

(AEOM-) AEOMICA INC.

New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.

Disclosure; SEQ ID NO 6213; 214pp; English

The present invention describes a human genome-derived myosin-like

C protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLPC l can be used in gene therapy and vaccine production. The hGDMLP-1

C nucleic acids can be used as probes to detect, characterise and quantify

nucleic acids can be used as probes to detect, characterise and quantify

C provide initial substrates for the recombinant engineering of hGDMLP-1

C protein variants having desired phenotypic improvements, and for

c protein variants having desired phenotypic improvements, and for

c protein variants having desired phenotypic improvements, and for

c sxpressing the proteins. The hGDMLP-1 proteins or polypeptides may be

cc used as immunogens to raise antibodies that specifically recognise hGDMLP

cused as immunogens to raise antibodies that specific blomclecule

and/or amount specifically of hGDMLP proteins, as specific blomclecule

capture probes for surface-enhanced laser desoribic of instantion, as

therefore associated with the expression of hGDMLP-1 in particular heart

c polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a

cc production, and in vaccines or for replacement therapy. The

production, and in vaccines or for replacement therapy. The

production, and in vaccines or for replacement therapy. The

computer associated with the expression of hGDMLP-1, in particular heart

can skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the

CT he present sequence data for this parent did not form part of the printed

c physicalication, but was obtained in electronic format directly from MIPO

c t ftp.wipo.int/pub/published_pct_esquence

Sequence 17 BP; 4 A; 7 C; 4 G; 2 T; 0 U; 0 Other;

Gaps ; 0 2.9%; Score 12.2; DB 1; Length 17; llarity 82.4%; Pred. No. 4.2e+02; Conservative 0; Mismatches 3; Indels Query Match Best Local Similarity Matches 14; Conserv

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 clambe used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser described in onisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                             Human, genome-derived myosin-like protein 1; GDMLP-1; hcDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:1014.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ji Y, Penn SG, Hanzel DK, Rank DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 1014; 214pp; English.
291 CTGGTGAAGGACCTGAG 307
                                                                                                                                ABN01022 standard; DNA; 17 BP.
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2000US-0236359P.
2000GB-00024263.
2001WO-US000661.
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30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000679.
50-JAN-2001; 2001WO-US000679.
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2001WO-US000663.
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                                                                                                                                                                                                         29-MAY-2002 (first entry)
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30-JAN-2001;
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Chen W, Shannon ME;

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New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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The present sequence represents an oligomer used in the screening of the hGDWLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; genome-derived myosin-like protein 1, GDMLP-1, hGDMLP-1, heart; muscle, myosin, chromosome 22, gene therapy, vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
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                                                                                                                                           Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                           Sequence 17 BP; 8 A; 1 C; 8 G; 0 T; 0 U; 0 Other;
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30-JNN-2001; 2001W0-US000661.
30-JNN-2001; 2001W0-US000663.
30-JNN-2001; 2001W0-US000663.
30-JNN-2001; 2001W0-US000664.
30-JNN-2001; 2001W0-US000665.
                                                                                                                                                                                                                     206 GAAAGCAGAGAACTCGG 222
                                                                                                                                                                                                                                                                                                                                              ABN00791 standard; DNA; 17 BP.
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30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
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27-SEP-2000;
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protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1 in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the chapture. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO. It the wipo.int/pub/published_pct_sequence
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Sequence 17 BP; 4 A; 6 C; 5 G; 2 T; 0 U; 0 Other;

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Gaps
Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                214 AGAACTCGGTGGCGCC 230
                                                                                                                                          17 AGATCTCGGTGCTGGCC 1
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Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:9021. ABN09029 standard; DNA; 17 BP 29-MAY-2002 (first entry) ABN09029; RESULT 768 ABN09029

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens.

WO200192524-A2.

36-DEC-2001,

25-MAY-2001; 2001WO-US016981 26-MAY-2000;

30-JAN-2001; 2001WO-US000663. 30-JAN-2001; 2001WO-US000664. 30-JAN-2001; 2001WO-US000665. 30-JAN-2001; 2001WO-US000666. 30-JAN-2001; 2001WO-US000666. 2000US-0234687P. 2000US-0236359P. 2000GB-00024263. 2001WO-US000661. 2001WO-US000662. 27-SEP-2000; 2 04-OCT-2000; 2 30-JAN-2001; 2 30-JAN-2001;

(AEOM-) AEOMICA INC.

30-JAN-2001; 2001WO-US000669

2001US-0266860P

Shannon ME; Chen W, Rank DR, Hanzel DK, Su Y, Ji Y, Penn SG, New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins,

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify conclude initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be capture amount specifically of hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specific blomclecule and/or amount specifically of hGDMLP-proteins, as specific blomclecule capture probes for surface-ahanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The concentration of skeleral muscle disorders. hGDMLP-1 may be used for diagnosing a cleorate associated with the expression of hGDMLP-1, in particular heart can skeleral muscle disorders. hGDMLP-1 is leaquence in the exemplification of the present invention. N.B. CT The sequence data for this patent did not form part of the printed capture. And the sequence of this patent did not form part of the printed capture properties of the sequence of the sequence.
or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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                                                                                             Disclosure, SEQ ID NO 9021; 214pp; English.
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Query Match
2.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 82.4%; Pred. No. 4.2e+02;
Matches 14; Conservative 0; Mismatches 3; Indels 289 AGCTGGTGAAGGACCTG 305

1 AGCTGGAGAAGTACGTG 17

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Gaps

Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:9919. ABN09927 standard; DNA; 17 BP. 29-MAY-2002 (first entry) ABN09927; RESULT 769 ABN09927/C

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens.

WO200192524-A2.

25-MAY-2001; 2001WO-US016981 26-MAY-2000; 2000US-0207456P 21-SEP-2000; 27-SEP-2000; 06-DEC-2001.

2001WO-US000663. 2001WO-US000664. 2001WO-US000665. 2001WO-US000666. 2001WO-US000667. 2001WO-US000668. 2001WO-US000661 2001WO-US000662 30-JAN-2001; 2 30-JAN-2001; 2 30-JAN-2001; 2 30-JAN-2001; 30-JAN-2001; 10-JAN-2001; Page 394

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Wed Apr 21 12:58:21 2004

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therappy and vaccine production. The hGDMLP-1 can be used as probes to detect, characterise and quantify nucleic acids in samples, as amplification substrates; to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser describing of inconsistion, as the present in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production and in vaccines or for replacement therapy. The production desequence represents an oligomer used in the screening of the hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                     New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                        Chen W,
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                                                                                                                                                      Rank DR,
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30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
05-FEB-2001; 2001US-0266860P.
                                                                                                                                                        Gu Y, Ji Y, Penn SG,
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                                                                                                     (AEOM-) AEOMICA INC
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New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.

Shannon ME;

Chen W,

Rank DR,

Hanzel DK,

Gu Y, Ji Y, Penn SG, (AEOM-) AEOMICA INC.

WPI; 2002-179446/23.

Shannon ME

2000US-0236359P

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Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels
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Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss. Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:1479. ABN01487 standard; DNA; 17 BP 71 CTACGAGGGCCGCGCAG 87 17 craagagagacrogoag 1 (first entry) 29-MAY-2002 ABN01487; RESULT 770 ABN01487/ à g RX PX SX XX XX BX PX XX BX XX

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The present invention describes a human genome-derived myosin-like protein I (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids in samples, as amplification substrates to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specific biomolecule and/or amount specifically of hGDMLP proteins, as specific biomolecule and/or amount specifically proteins, as specific biomolecule capture probes for surface-enhanced laser describing in onisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the home represents an oligomer used in the screening of the home how the present invention. N.B. The sequence data for this patent did not form part of the printed of the present invention. N.B. The print of the present invention.
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Human, KTOM1a; KTOM1; kidney tumour overexpressed membrane; cytostatic; gene therapy; cancer; kidney; liver; bone marrow; brain; heart; lung;

25-MAY-2001; 2001WO-US016981 26-MAY-2000; 2000US-0207456P.

WO200192524-A2

06-DEC-2001

Homo sapiens.

kidney; colon; skeletal muscle; testis; uterus; placenta; probe; ss.

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gene therapy; cancer; kidney; liver; bone marrow; brain; heart; lung;
kidney; colon; skeletal muscle; testis; uterus; placenta; probe; ss.
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27-SEP-2000; 2000US-0235559P.
04-OCT-2001; 2001WG-02004663.
30-JAN-2001; 2001WG-US000663.
30-JAN-2001; 2001WG-US000663.
30-JAN-2001; 2001WG-US0006663.
30-JAN-2001; 2001WG-US000666.
30-JAN-2001; 2001WG-US000666.
30-JAN-2001; 2001WG-US000667.
30-JAN-2001; 2001WG-US000669.
30-JAN-2001; 2001WG-US000669.
30-JAN-2001; 2001WG-US000669.
23-WAY-2001; 2001WG-US000669.
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Best Local Similarity 82.4%;
Matches 14; Conservative
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                                                           Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New human kidney tumor overexpressed membrane (KTOM1) protein and nucleic acids encoding the protein, useful for treating subjects having defects in KTOM1 which can manifest as cancer of the kidney, or as a disorder of e.g., liver or bone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a novel isolated nucleic acid encoding human KTOM1 (kidney tumour overexpressed membrane) protein. The protein of the threethor has cyrostatic activity. The nucleotide may have a use in gene therapy. The KTOM1 nucleic acids may be used to diagnose, treat or monitor a disease caused by altered expression of human KTOM1 compositions comprising the nucleic acids in KTOM1 which can manifest as used to treat subjects having defects in KTOM1 which can manifest as cancer of the kidney, as well as a disorder of liver, bone marrow, brain, heart, lung, kidney, only, skeletal muscle, restis, uterus and placenta function. The sequence represents a probe used in the invention to scan the nt 1-1001 portion of human KTOM1a (ABQ63232)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 2; Page 165; 418pp; English.
                                                                                                                                                                                 21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0236359P.
04-OCT-2000; 2000US-0236359P.
30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000667.
23-MAY-2001; 2001WO-US000677.
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                                                                                                                                                 21-SEP-2001; 2001WO-US029656
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    43 ATGGCCACCACTCAGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABQ63351 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-AUG-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-479509/51
                                                                        WO200224750-A2
                                       Homo sapiens.
                                                                                                              28-MAR-2002
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Shang J;

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New human kidney tumor overexpressed membrane (KTOM1) protein and nucleic acids encoding the protein, useful for treating subjects having defects in KTOM1 which can manifest as cancer of the kidney, or as a disorder of e.g., liver or bone.
                                                                                                                                                                                                                                    The invention relates to a novel isolated nucleic acid encoding human KTOM1 (kidney tumour overexpressed membrane) protein. The protein of the throention has cyrostatic activity. The nucleotide may have a use in gene therapy. The KTOM1 nucleic acids may be used to diagnose, treat or monitor a disease caused by altered expression of human KTOM1. Compositions comprising the nucleic acids, proteins or antibodies may be used to treat subjects having defects in KTOM1 which can manifest as cancer of the kidney, as well as a disorder of liver, bone marrow, brain, heart, lung, kidney, colon, skeletal muscle, testis, uterus and placenta function. The sequence represents a probe used in the invention to scan the nt 1-1001 portion of human KTOM1a (ABQ63123)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17 BP; 4 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                               Example 2; Page 166; 418pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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ABV85548
XX
XX
AC ABV8
XX
DT 11-D
XX
XX
XX
XX
XX
XX
XX
XX
XX
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Human; KTOMla; KTOMl; kidney tumour overexpressed membrane; cytostatic;

Human KTOM1a portion (ABQ63232) probe # 64.

ABQ63351;

ABQ63351 ID ABQ XX AC ABQ XX DT 20-XX DE HUM XX KW HUM

RESULT 772

à g

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Homo sapiens.
Synthetic.
                                              EP1243660-A2
 The present invention describes an isolated nucleic acid (I) encoding a human UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase 10 (EpadNarase 10, EC 24.1.41) protein. Human pp-GaNTase 10 is located to chromosome 7q11.2. (I) can be used in gene therapy. Molecules of the present invention can be used in therapy, particularly to prevent or treat a disorder associated with decreased expression or activity of EpadNarase. The sequences given in ABV85011 to ABV8668 and ABP53502 to ABP53504 are given in the exemplification of the present invention. N.B. The sequence data for this patent is not represented in the printed specification but is based on sequence information supplied by the
                                                                                                                                                                                                                                                                                                                                                 Nucleic acid encoding human UDP-GalNAc:polypeptide N-cetylgalactosaminyltransferase 10 protein is useful to diagnose, prevent and treat disorders associated with reduced or over expression of the
Human, UDP-GalNAc.polypeptide N-acetylgalactosaminyltransferase 10;
pp-GaNTase 10; EC 2.4.1.41; chromosome 7q11.2; gene therapy; scanning;
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ilarity 82.4%; Pred. No. 4.2e+02;
Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 4 A; 4 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; SEQ ID NO 541; 59pp; English.
                                                                                                                                                           30-JNN-2001; 2001WO-US000664.
30-JNN-2001; 2001WO-US000665.
30-JNN-2001; 2001WO-US000666.
30-JNN-2001; 2001WO-US000667.
30-JNN-2001; 2001WO-US000669.
30-JNN-2001; 2001WO-US000669.
23-MAY-2001; 2001WS-US000669.
30-JNN-2001; 2001WS-US000677.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         403 TCTTCTACGTGATCGAG 419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABV85708 standard; DNA; 17 BP.
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                                                                                                                             25-JAN-2002; 2002EP-00001161
                                                                                                                                                                                                                                                                                                    Zhang J, Gu Y, Nguyen C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
es 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                  encoded protein.
                                              Homo sapiens.
Synthetic.
                                                                               EP1243660-A2
                                                                                                      25-SEP-2002
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IID ABV8
XX
AC ABV8
XX
DT 11-D
XX
DE Huma
XX
KW Huma
KW SS.
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Nucleic acid encoding human UDP-GalNAc:polypeptide N-cetylgalactosaminyltransferase 10 protein is useful to diagnose, prevent and treat disorders associated with reduced or over expression of the encoded protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes an isolated nucleic acid (I) encoding a human UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase 10 (pp-daNTase 10, EC 2.4.1.41) protein. Human pp-daNTase 10 is located to chromosome 7q11.2. (I) can be used in gene therapy. Molecules of the present invention can be used in therapy, particularly to prevent or treat a disorder associated with decreased expression or activity of pp-daNTase. The sequences given in ABV85011 to ABV86689 and ABP53502 to ABP53504 are given in the exemplification of the present invention. N.B. The sequence data for this patent is not represented in the printed specification but is based on sequence information supplied by the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human, gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; afteral; liver; male germ cell development; bone marrow; brain; kidhey; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 2 A; 5 C; 8 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; SEQ ID NO 701; 59pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     284 CACCAAGCTGGTGAAGG 300
                                                                                                                                     30-JAN-2001; 2001WC-US000666.
30-JAN-2001; 2001WC-US000667.
30-JAN-2001; 2001WC-US000668.
30-JAN-2001; 2001WC-US000669.
30-JAN-2001; 2001WC-US0006770.
23-XAX-2001; 2001US-00864761.
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Zhang J, Gu Y, Nguyen C;
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                                                                                                                                                                                                                                                                                                                                                                      (AEOM-) AEOMICA INC.
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                                                    30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
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25-JAN-2002;
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ABV79551/c
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Human, UDP-GalNAc.polypeptide N-acetylgalactosaminyltransferase 10; pp-GaNTase 10; EC 2.4.1.41; chromosome 7q11.2; gene therapy; scanning; ss.

Human pp-GaNTase 10 scanning 17-mer SEQ ID NO:701.

11-DEC-2002 (first entry)

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Wed Apr 21 12:58:21 2004
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Homo sapiens EP1239051-A2

rng.res

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The present invention relates to human testis expressed Patched like protein (HTPL, see ABV78759 to ABV78762 and ABB9819 to ABB98520). HTPL has two isoforms, with a few single base pair differences between the two of the single base pair differences between the two of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL shares an overall structure organisation with the Patched protein. The shares an overall structure organisation with the Patched protein. The shares an overall structure organisation with the Patched protein. The shared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene was important in regulating male germ cell development, and the HTPL, and in therapy and manufacture of a medicament for treatment or prevention of therapy and manufacture of a medicament for treatment or prevention of the HTPL. Such disorders include disorders expression or activity of human HTPL. Such disorders include disorders of testis, or adrenal, adult and foctal liver, bone marrow, brain, kidney, lung, placenta, prostate, skeletal muscle or colon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potenial therapeutic agents for male infertility and cancer. The present oligonucleotide was used in an example from the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; Page 168; 718pp; English.
                                                                                                                                                              30-JAN-2001, 2001WO-US000663.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000668.
33-JAN-2001; 2001WO-US000669.
23-MAX-2001; 2001WS-0800669.
09-OCT-2001; 2001US-0300669.
                                                                                                                     28-JAN-2002; 2002EP-00001167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-676582/73.
                                                                                                                                                                                                                                                                                                                                                                                    (AEOM-) AEOMICA INC
                                                                       07-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                    Zhan J;
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0; Gaps Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels Sequence 17 BP; 3 A; 4 C; 9 G; 1 T; 0 U; 0 Other;

373 TCCTGGACCGCGACGAC 389

Human POSHL1 scanning oligonucleotide SEQ ID NO 1746.

Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine;

30-JAN-2001, 2001WO-US000665. 30-JAN-2001, 2001WO-US000666. 30-JAN-2001, 2001WO-US000667. 30-JAN-2001, 2001WO-US000667. 30-JAN-2001, 2001WO-US000669. 30-JAN-2001, 2001WG-US0006670. 23-WAX-2001, 2001US-00864761. 2002EP-00001165 WPI; 2002-684061/74. (AEOM-) AEOMICA INC. 28-JAN-2002; 30-JAN-2001; 11-SEP-2002. Shannon M;

The invention relates to an isolated SH3 domain (POSH)-like signalling cortein 1 (POSHI 1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABBB3939), a sequence having 65% sequence identity to (S1), (S1), having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHI is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPsases as well as downstream components of the signal transduction pathway. (I) is useful of contentifying a specific binding partner. (I) and nucleic acids (II) for identifying a specific binding partner. (I) and nucleic acids (II) consecting (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHI including diagnosing and crusting caused by altered expression of human POSHI including microarrays which are useful for measuring and for surveying gene expression and creating cransgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligomodicotide useful in examples cof the invention. Note: The present sequence did not form part of the content by the European Patent Office Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1. Example 2; SEQ ID NO 1746; 60pp + Sequence Listing; English

0; Gaps Query Match
2.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 82.4%; Pred. No. 4.2e+02;
Matches 14; Conservative 0; Mismatches 3; Indels Seguence 17 BP; 4 A; 7 C; 5 G; 1 T; 0 U; 0 Other;

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ABL31783 standard; DNA; 17 BP

ABL31783/c

ABL31783;

Human HLA genotyping oligonucleotide SEQ ID NO 1272. 21-MAR-2002 (first entry)

Human; human leukocyte antigen; HLA; genotype; polymorphism;

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The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligonucleotides (AB130512-AB131809) originating in the sequences of genes (AB130512-AB131809) originating in the sequences of pene containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals
                                                                                                                                                                                                                                                                                                                                                                     Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when transplanting between them.
                                                                                                                                                                                                                                                                                             Ichihara T, Matsumura Y, Moriya S, Nishida M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ch 2.9%; Score 12.2; DB 1; Length 17; 1 Similarity 82.4%; Pred. No. 4.2e+02; 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 3 A; 7 C; 6 G; 1 T; 0 U; 0 Other;
immunogenetic; transplantation; genetic disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 10; Page 334; 345pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             304 TGAGCCCCGGGGACCGC 320
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                                                                                                                                                                                                01-JUN-2000; 2000JP-00164798.
                                                                                                                                                        01-JUN-2001; 2001WO-JP004662
                                                                                                                                                                                                                                    (NISN ) NISSHINBO IND INC
(SYST-) SYSTEM RES INC.
                                                                                                                                                                                                                                                                                               noko H, Kagiya T,
                                                                                                                                                                                                                                                                                                                                   WPI; 2002-122074/16.
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                                                                              WO200192572-A1
                                          Homo sapiens.
                                                                                                                    06-DEC-2001
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Matches
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0; Gaps
                                                                                                     ABK56639 standard; RNA; 17 BP
                                                                                                                                                   (first entry)
                                                                                                                                                   02-JUL-2002
                                                                                                                              ABK56639;
                                                                               RESULT 778
                                                                                             ABK56639/c
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BP

ABZ95537 standard; DNA; 17

(first entry)

17-OCT-2003

ABZ95537;

Human; chloride channel calcium activated 1; CLCA1; ss; antiasthmatic; antinflammatory; chronic obstructive pulmonary disease; COPD; asthma; chronic bronchitis; cystic fibrosis; obstructive bowel syndrome; oxygen therapy; bronchodilator; corticosteroid; vaccination; mucokinetic; acetylcysteine

Human CLCAl gene enzymatic nucleic acid #1010.

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09-AUG-2001; 2001WO-US024970.
                                           WO200211674-A2
                                      Ното варіепв.
                                                 14-FEB-2002
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The invention relates to enzymatic nucleic acid molecules that down of regulate expression of chloride channel calcium activated 1 (CLCA1) genes by cleaving RNA derived from the genes. The nucleic acid sequences are cuseful as pharmaceutical agents for treating conditions such as chronic obstructive pulmonary disease (CPD), chronic bronchitis, asthma, cystic fibrosis, obstructive bowel syndrome and any other diseases or conditions that are related to or will respond to the levels of CLCA1 in a cell or tissue. The sequences are useful for reducing CLCA1 activity in a cell, chence, are useful for treatment of a patient having a condition associated with the level of CLCA1, where the invention further comprises treatment, for example, oxygen therapies under conditions suitable for the treatment, for example, oxygen therapy, bronchodilators, corticosteroids, antibacterials, vaccinations, acetylcysteine and mucoshinetic agents. The nucleic acids of the invention are also used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of CLCA1 RNA in a cell. This sequence represents an
                                                                                                                                                                                                                                              Enzymatic polynucleotide that down regulates expression of chloride channel calcium activated gene, useful for treating Chronic obstructive pulmonary disease (COPD), chronic bronchitis and asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                Thompson J, Mcswiggen J, Mckenzie T, Ayers D, Szymkowski DB;
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2.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 82.4%; Pred. No. 4.2e+02;
Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 5 A; 6 C; 3 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          enzymatic nucleic acid molecule of the invention
                                                                                                                                                                                                                                                                                                                                                     Claim 4; Page 76; 152pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      231 AAATCGGGAGGCTGCTT 247
09-AUG-2000; 2000US-0224383P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17 AATTGGGGAGGCTCCTT
                                             (RIBO-) RIBOZYME PHARM INC.
                                                                        USA LLC
                                                                                                                                                                                                           WPI; 2002-217145/27.
                                                                   (SYNT ) SYNTEX USA (THOM/) THOMPSON J.
                                                                                                                                                                   Grupe A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 779
ABZ95537/c
ID ABZ9553
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Human, antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. Human endothelial nitric oxide synthase antisense fragment no.1401.

23-APR-2002; 2002WO-US013135. 24-APR-2001; 2001US-0286137P Homo sapiens 31-OCT-2002

```
The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions; 5' and 3' intron-exon junctions, or regions within 2-10 mucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and associated with lung and/or antiinflammatory steroid and ubiquinone. A composition of the invention and antiinflammatory, antiallergic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, as 1so for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of of acceptor, producing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing bronchoconstriction, lung allergies, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition.

Unug inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                   Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; SEQ ID NO 10779; 872pp; English.
                                                                                                                 Katz E,
                                                                                                     Li Y, Sandrasagra A, K
Tang L, Shahabuddin S;
(EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                   WPI; 2003-229219/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ubiquinone
                                                                                                                                                                 Miller S,
                                                                                                           Nyce JW,
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ö 2.9%; Score 12.2; DB 1; Length 17; larity 82.4%; Pred, No. 4.2e+02; Conservative 0; Mismatches 3; Indels Sequence 17 BP; 0 A; 7 C; 6 G; 4 T; 0 U; 0 Other; ftp.wipo.int/pub/published_pct_sequences Similarity 14; Query Match Best Local Matches

268 ACCTGGAGCAGGGGGGC 284 17 Accedeadeadeaced 1 셤 ઠે

Human PDE4A-MTA oligonucleotide sequence. ABZ99035 standard; DNA; 17 BP (first entry) 17-0CT-2003 ABZ99035; RESULT 780 ABZ99035

Human, antisense, lung dysfunction, nasal airway dysfunction, antinflammatory steroid, ubiquinone, antinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive, cytostatic; gene therapy, antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

#0200285308-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Aguilar D;

Pabalan J,

Aguilar D; Katz E, Pabalan J, Li Y, Sandrasagra A, Tang L, Shahabuddin Nyce JW,] Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 14277; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antisense to the intitation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 muclectides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antisthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a tenential a natistense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing levels of or reducing sensitivity to adenosine, reducing levels of or reducing sensitivity to adenosine, reducing levels of ung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence date for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO ftp.wipo.int/pub/published_pct_sequences

Seguence 17 BP; 3 A; 6 C; 5 G; 3 T; 0 U; 0 Other;

Gaps ö Query Match
2.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 82.4%; Pred. No. 4.2e+02;
Matches 14; Conservative 0; Mismatches 3; Indels

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Gaps

Lactobacillus brevis PCR primer ORF4 SEQ ID NO:66. (first entry) 29-APR-2003 ABZ76563;

ABZ76563 standard; DNA; 17 BP.

RESULT 781 ABZ76563

Lactobacillus brevis; beer turbidity; beer clouding; beer; detection; lactic acid bacteria; brewing; probe; PCR primer; ss. XLXBXBXBXBXBXBXBXBXBXBXBXBXX

Lactobacillus brevis.

WO200295028-A1.

28-NOV-2002.

23-MAY-2002; 2002WO-JP005022.

23-MAY-2001; 2001JP-00154085.

(KIRI) KIRIN BEER KK.

Ë Fujii ö

Gaps

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WPI; 2003-250498/25.
WPI; 2003-120803/11
                                                     Local Similarity
                                                                                                           Homo sapiens
                                                                                                                FR2826373-A1
                                                                                                                                        ruijnder M,
                                                                                       27-JUN-2003
                                                                                                                     27-DEC-2002
                                         invention
                                                                                   ACC51810;
                                                   Query Match
                                                                         RESULT 782
                                                        Matches
                                                                           ACC51810
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The invention describes an isolated nucleic acid encoding a G protein coupled receptor (GPCR), mutations of which cause cancer, comprising a 2225 or 1921 base pair sequence, or their degenerate variants, encoding a 409 residue amino acid sequence, all given in the specification, with or without conservative amino acid substitutions, or complements of the sequence of them. The encoding nucleic acid is not more than 100 kbase in length. The methods and compositions of the present invention are useful for diagnosing, investigating and/or treating disorders associated with aberrant expression or activity of GPCR-A-1, such as tumours and cancers. This sequence represents an oligonucleotide used to analyse the gene encoding human G-protein coupled receptor GPCR-A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human, G-protein coupled receptor; GPCR-A-1; cancer; tumour;
G-Protein-Agonist; G-Protein-Antagonist; gene therapy; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   G-protein coupled receptor GPCR-A-1 analysis oligonucleotide #187.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New GPCR-A-1 nucleic acid and polypeptide, useful for diagnosing, investigating and/or treating disorders associated with aberrant expression or activity of GPCR-A-1, such as tumors and cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.9%; Score 12.2; DB 1; Length 17; 12.4%; Pred. No. 4.2e+02; ve 0; Mismatches 3; Indels
                                                                2.9%; Score 12.2; DB 1; Length 17;
ilarity 82.4%; Pred. No. 4.2e+02;
Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 4 A; 3 C; 7 G; 3 T; 0 U; 0 Other;
                  Seguence 17 BP; 6 A; 4 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 2; SEQ ID NO 211; 156pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (AMSH ) AMERSHAM BIOSCIENCES SV CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   304
                                                                                                                                                                         289
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                                                                                                                                                                                                                                                                                                                                                     ACA99694 standard; DNA; 17 BP.
                                                                                                                                                                                                                    1 darcadddddddau 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-OCT-2001; 2001US-0329000P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           82.48;
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ABT37105/c
ID ABT37105 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      288 AAGCTGGTGAAGGACCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 AAGCTGGTAGGGGACCT
                                                                                                                                                                      273 GAGCAGGCGGCACCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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ses 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-381720/36.
                                                                                           Local Similarity
es 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003031621-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                     ACA99694;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Zhang J;
                                                                        Query Match
                                                                                                                                                                                                                                                                                                    RESULT 783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                            The present invention describes a polynucleotide probe, or primer, for detecting beer-clouding lactic acid bacteria containing a nucleotide sequence of (1) with 8056 base pairs (see ABZ/6501), or a nucleotide made from not less than 15 nucleotides hybridisable with its complementary sequence. Probes and primers from the present invention can be used for detecting beer-clouding lactic acid bacteria (Lactobacillus brevis) for quality control during beer production, which is applicable in the brewing industry. The present sequence represents a PCR primer for lactobacillus brevis which is used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New nucleic acid sequences associated with tumor suppression, regression, apoptosis or virus resistance are useful to diagnose and treat viral disease, development of tumor cells and cell degeneration.
                                             Polynucleotide probes and primers for detecting beer-clouding lactic acid bacteria, for quality control during beer production applicable in brewing industry.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      with tundur suppression or regression, apoptosis or virus resistance. invention relates to these sequences or sequences having at least 80% identity to them, and polypeptides encoded by the sequences or polypeptides having 80% identity to the polypeptide sequences. The invention is used to diagnose or treat viral disease or disease characterized by development of tumour cells or cellular degeneration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This sequence represents an isolated nucleic acid sequence associated with tumour suppression or regression, apoptosis or virus resistance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ss; tumour suppressor; antitumour; cytostatic; tumour suppression; tumour regression; apoptosis; virus resistance; diagnosis; cellular degeneration.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 4.2e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 5 A; 4 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human tumour suppressor sequence #577.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Amson R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 173; 798pp; French
                                                                                                                                                   Claim 7; Page 31; 94pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       225 GCGGCCAAATCGGGAGG 241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACC51810 standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-JUN-2001; 2001FR-00008139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 gcagccaaarcgrearg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MOLE-) MOLECULAR ENGINES
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Gaps

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(first entry)

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(MOLE-) MOLECULAR ENGINES LAB.
                                    felerman A, Amson R,
                                      WPI; 2003-313353/30.
    12-JUN-2003
 ABT37105;
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Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.
Tumour suppression related human fukutin oligo SEQ ID No 2742.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-SEP-2001; 2001FR-00011978.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-SEP-2002; 2002WO-IB004208
                                                                                                                                                                                                                                                                                                                                    WO2003025175-A2.
                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                            27-MAR-2003.
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New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.

Tuijnder M;

Disclosure, Page 353; 720pp; French

The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that tybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated mucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti) sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, production of paramaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell diseases that are characterised by development of tumours or cell diseases that are characterised by development of tumours or cell diseases that are characterised by development of tumours or cell diseases. The polypeptides can also be used to generate antibodies, and cher the polypeptides can also be used to generate antibodies, and cher the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene therapy. This polymuclectide sequence represents a tumour suppression characterial at the invention can be used in gene thorap.

Sequence 17 BP; 2 A; 7 C; 2 G; 6 T; 0 U; 0 Other;

Gaps . 0 2.9%; Score 12.2; DB 1; Length 17; Local Similarity 82.4%; Pred. No. 4.26+02; les 14; Conservative 0; Mismarchar

287 CAAGCTGGTGAAGGACC 303 chadgaddrichaddarc 1

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RESULT 785 ABT34651/c ID ABT34651 standard; DNA; 17 BP. XX AC ABT34651:

12-JUN-2003 (first entry)

Tumour suppression related human fukutin oligo SEQ ID No 288

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.

Homo sapiens

WO2003025175-A2.

27-MAR-2003

17-SEP-2002; 2002WO-IB004208

17-SEP-2001; 2001FR-00011978

(MOLE-) MOLECULAR ENGINES LAB.

Tuijnder Amson R, relerman A,

Σ

WPI; 2003-313353/30

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.

Disclosure; Page 67; 720pp; French.

The invention relates to a novel isolated 17 mer nucleic acid sequence, can given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal all alignment, at least 80 % identity to the 17 mer sequence, a sequence that bybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated mucleic acids of the invention are useful as probes and primers for detecting, acids of the invention are useful as probes and primers for detecting, in vitro as (anti) sense reagents, and for production of recombinant polypeptides and primers for detecting, production of recombinant polypeptides. Any of the nucleic acids, propublicies, vectors containing the nucleic acids, colls containing the nucleic acids, cells containing the operation of plarmaceuticials for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell diseases that are characterised by development of tumours or cell diseases that any properties can also be used to generate antibodies, and contain the polypeptides can also be used to generate antibodies, and chops. The nucleic acid sequences of the invention can be used in gene therapy. This polynucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention

Seguence 17 BP; 3 A; 1 C; 8 G; 5 T; 0 U; 0 Other;

Gaps ö 2.9%; Score 12.2; DB 1; Length 17; 32.4%; Pred. No. 4.2e+02; ive 0; Mismatches 3; Indels Local Similarity 82.4%; hes 14; Conservative Query Match Best Loca Matches

92 CATCACCACGTCTGACC 108

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17 CAACACCACCTCTGATC 1464/c ABT37464 standard; DNA; 17 RESULT 786 ABT37464/

ABT37464;

(first entry) 12-JUN-2003

Tumour suppression related human fukutin oligo SEQ ID No 3101

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; attisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.

Homo sapiens

402003025175-A2

17-SEP-2002; 2002WO-IB004208

17-SEP-2001; 2001FR-00011978

Tuijnder M; relerman A, Amson R,

WPI; 2003-313353/30.

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.

The invention relates to a novel isolated 17 mer mucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence with, after optimal alloguent, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated mucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, e.g. as one caids of the invention are useful sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, colypeptides, vectors containing the nucleic acids, colypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell diseases that are characterised by development of tumours or cell diseases that are characterised by development of tumours or tell patient samples is useful for disapnosis and/or prognosis of these containing the polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene thorapy. This polymuclectide sequence represents a tumour suppression characy. This polymuclectide sequence represents a tumour suppression

Sequence 17 BP; 2 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

g

03-JUN-2003

G-cleaver, amberzyme, cancer, REL-A activity, breast cancer; human, lung cancer; prostate cancer; colorectal cancer; brain cancer; hosead cancer; broaded cancer; brain cancer; cervical cancer; cancer; covarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; displatin; methotrexate; cyclophosphamide; docetaxel; displatin; methotrexate; centralitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic alrway inflammatory bowel disease; infection; ss. Bnzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme;

Homo sapiens.

US2002177568-A1.

28-NOV-2002.

23-MAY-2001; 2001US-00864785

92US-009B7132. 94US-00245466. 94US-00291932. 96US-00777916. 18-MAY-1994;

23-DEC-1996;

(STIN/) STINCHCOMB D (MCSW/) MCSWIGGEN J. (MCSW/) MCSWIGGEN J. (DRAP/) DRAPER K G. Stinchcomb DT, Mcswiggen J, Draper KG;

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 41; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFKA), where (I) is an inozyme, zinzwe, q-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg^27+. The enzymatic and antiense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, ossophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multiputor sesistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, datexaxte, cyclophosphamide, doxorubin, fluorouracil carboplatin, datexaxte, cyclophosphamide, doxorubin, fluorouracil carboplatin, datesate, cyclophosphamide, doxorubin, fluorouracil carboplatin, datesate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatexaxte, colocitio, gene therapy applications, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CMS) and myocardial), glomerulomes or a infection. This sequence represents the subs

Sequence 17 BP; 2 A; 6 C; 6 G; 0 T; 3 U; 0 Other;

Gaps ; 0 Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 70.6%; Pred. No. 4.2e+02; Matches 12; Conservative 2; Mismatches 3; Indels

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(MOLE-) MOLECULAR ENGINES LAB.

Disclosure; Page 395; 720pp; French.

Gaps ö Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels

CCTGAGCCCCGGGGACC 318 ccreacccaeccanc 17

ACA07885 standard; RNA; 17 BP RESULT 787 ACA07885 BXBXBXB

NFKB sub-unit modulating zinzyme substrate #284.

(first entry)

G-cleaver, amberzyment, moretem activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; cervical cancer; pancreatic cancer; cervical cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemocherapy; pacliteat; docetaxel; cipplatin; methotrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edarexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenceis; Crohn's disease; obeelty; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammatory bowel disease; infection; ss. Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; NFKB sub-unit modulating inozyme substrate #263. ACA06444 standard; RNA; 17 BP. (first entry) 03-JUN-2003 ACA06444;

US2002177568-A1. Homo sapiens.

28-NOV-2002.

92US-00987132. 94US-00245466. 94US-00291932. 96US-00777916. 23-MAY-2001; 2001US-00864785. 18-MAY-1994; 15-AUG-1994; 23-DEC-1996; 17-DEC-1992;

STINCHCOMB D T. MCSWIGGEN J. DRAPER K G. (MCSW/) N (DRAP/) I (STIN/)

Mcswiggen J, Stinchcomb DT,

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 31; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFRM), where (I) is an inozyme, zinzyme, q-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg^2, The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oeophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, gencitabine or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as

rheumatoid arthritis, restenosis, asthma, Crohn's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, alsepsis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic nucleic acid molecule 8888888888

Sequence 17 BP; 2 A; 9 C; 4 G; 0 T; 2 U; 0 Other;

ö Query Match
2.9%; Score 12.2; DB 1; Length 17;
Best Local Similarity 82.4%; Pred. No. 4.2e+02;
Matches 14; Conservative 0; Mismatches 3; Indels

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ACA06721 standard; RNA; 17 BP. ACA06721,

ACA06721;

03-JUN-2003 (first entry)

NFKB sub-unit modulating inozyme substrate #540.

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; dung cancer; brain cancer; brostete cancer; colorectal cancer; brain cancer; cesophageal cancer; stomach cancer; bladder cancer; pancreatic cancer; cervical cancer; head and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A specific inhibitor; confontherapy; paclitaxel; docetaxel; cisplatin; methorraxate; confontherapy; acclitaxel; duocetaxel; cisplatin; methorraxate; gencitabine; radiation therapy; inflammatory diseas; asthma; diabetes; gencitabine; radiation therapy; inflammatory diseas; asthma; diabetes; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; ss.

Homo sapiens.

U\$2002177568-A1.

28-NOV-2002.

23-MAY-2001; 2001US-00864785

92US-00987132. 94US-00245466. 94US-00291932. 96US-00777916. 18-MAY-1994; 23-DEC-1996; 07-DEC-1992;

(STIN/) STINCHCOMB D T. (MCSW/) MCSWIGGEN J (MCSW/) MCSWIGGEN J. (DRAP/) DRAPER K G. Draper KG; Mcswiggen J, Stinchcomb DT,

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 35; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFKB), where (I) is an inozyme, zinzyme, G-cleaver or amberzyme

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countigutation, the enapymention associated with the level of treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A.

(I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG-2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophagel, stoomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specialic inhibitors or chemotherapy including paclitaxal, docetaxel, cisplatin, methotrexate, gemiciabine or radiation fluorouracil carboplatin, edatrexate, gemiciabine or radiation therapy. The enzymatic and antisense nucleic coid molecules are also useful for treating inflammatory disease such as theumatoid arthritis, restenosis, asthma, Crohm's disease, diabetes, cheumatoid arthritis, restenosis, used scleaving inflammatory disease, caid asterial nervous system (CNS) and myocardial), glomerulonephritis, septicin, myis conversed to a novel pervonment of a novel pervonment of a novel pervonment.
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configuration. The enzymatic nucleic acid molecule is adapted to treat
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Sequence 17 BP; 2 A; 3 C; 5 G; 0 T; 7 U; 0 Other;

Gaps ö Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels

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ACA09012 standard; RNA; 17 BP. ACA09012; RESULT 790 ACA09012

03-JUN-2003 (first entry)

NFKB sub-unit modulating amberzyme substrate #175.

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; cesophageal cancer; stomach cancer; bladder cancer; pancreatic cancer; cervical cancer; head and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; commontarpy; paclitexap; paplatin; methorrexate; colomotherapy; paclitexap; docetaxel; cisplatin; methorrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatorid arthritis; restences; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfushon injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; ss.

Homo sapiens.

US2002177568-A1.

28-NOV-2002.

23-MAY-2001; 2001US-00864785.

92US-00987132. 94US-00245466. 94US-00291932. 96US-00777916. 07-DEC-1992; 18-MAY-1994;

STINCHCOMB D T. MCSWIGGEN J. DRAPER K G. 33-DEC-1996; (STIN/) (MCSW/)

Draper KG Mcswiggen J, Stinchcomb DT,

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 54; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFKB), where (I) is an inozyme, ginzyme, g-clasver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG^2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or cervical, essistant cancer. The method involves use of other drug trempted therapy including pacilitaxel, docetaxel, displath, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplath, edarrexate, cyclophosphamide, doxorubin, fluorouracil carboplath, edarrexate, cyclophosphamide, doxorubin, fluorouracil carboplath, edarrexate, commatoid arthritis, restenosis, asthma, Crohn's disease such as cheening and seases, lupus, multiple sclerosis, transplant/graft cirval interval inflammatory disease, lupus, multiple sclerosis, transplant/graft cirval and encourant perpanents he annowed represented in pervone system (CNS) and myocardial), glomerulonephritis, especies, allergic airvay inflammatory annowed disease or inferrion, and sease, allergic airvay inflammatory annowed disease or inferrion, and encourant pervones represented her annowed encouragemented annowed disease or inferrion, and encouragemented heromy annowed disease or inferrion, and encouragemented heromy annowed disease or inferrion, and encouragemented heromy and encouragemented heromy and encouragemented heromy and encouragemented infection. This sequence represents the substrate of a novel enzymatic mucleic acid molecule

Sequence 17 BP; 5 A; 3 C; 8 G; 0 T; 1 U; 0 Other;

0; Gaps 2.9%; Score 12.2; DB 1; Length 17; 76.5%; Pred. No. 4.2e+02; Live 1; Mismatches 3; Indel8 Best Local Similarity 76.5 Matches 13; Conservative Query Match

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ADA99253 standard; DNA; 17 BP ADA99253; RESULT 791 ADA99253/c

20-NOV-2003 (first entry)

Human MDZ3 scanning oligonucleotide SEQ ID 242.

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MD23; MD24; MD27; MD212; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

Homo sapiens.

EP1281758-A2.

35-FEB-2003.

30-JUL-2002; 2002EP-00016874.

02-AUG-2001; 2001US-00922181.

(AEOM-) AEOMICA INC

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Gu Y, Nguyen C;
                        WPI; 2003-423107/40
Shannon M,
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New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 242; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD24, MD24, MD212. MD23 is proteins and their coding sequences: MD24, MD27, MD212. MD23 is encoded at chromosome 7621.1, MD24 is encoded at chromosome 6921.3-22.2, MD27 is encoded at chromosome 6921.3-22.2, MD27 is encoded at chromosome 15q26.1. The MD23, MD24, MD27, and MD212 sequences are useful in therapy, or in manifacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23, MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as proteins. The present sequence was used to illustrate the invention.

Sequence 17 BP; 4 A; 4 C; 3 G; 6 T; 0 U; 0 Other;

Gaps ö Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels

57 GAGGAGTCTCTGCACTA 73

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RESULT 792 ADA99417

ADA99417 standard; DNA; 17 BP ADA99417;

(first entry)

20-NOV-2003

Human MDZ3 scanning oligonucleotide SEQ ID 406.

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

3P1281758-A2.

05-FEB-2003;

30-JUL-2002; 2002EP-00016874

02-AUG-2001; 2001US-00922181.

(AEOM-) AEOMICA INC.

Shannon M, Gu Y, Nguyen C;

WPI; 2003-423107/40.

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3,

MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 406; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ12. MDZ3 is encoded at chromosome 7422.1. MDZ4 is encoded at chromosome 6721.3-22.2.

MDZ7 is encoded at chromosome 16011.2 and MDZ12 is encoded at chromosome contains an edicament for treating or preventing at chromosome of polymers, and MDZ12 sequences are useful in therapy, or immunifacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross acids can also be used as probes to detect and characterize gross catterations in MDZ3, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as proteins are useful as therapeutic agents for gene therapy or as

Sequence 17 BP; 2 A; 9 C; 2 G; 4 T; 0 U; 0 Other;

2.9%; Score 12.2; DB 1; Length 17; 32.4%; Pred. No. 4.2e+02; ved. 0; Mismatches 3; Indels Local Similarity 82.4%; nes 14; Conservative Query Match Best Loca Matches

RESULT 793 ADB00316

ADB00316 standard; DNA; 17 BP

ADB00316;

20-NOV-2003 (first entry)

Human MDZ3 scanning oligonucleotide SEQ ID 1302.

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; se.

Homo sapiens

EP1281758-A2.

05-FEB-2003.

30-JUL-2002; 2002EP-00016874

02-AUG-2001; 2001US-00922181.

(AEOM-) AEOMICA INC.

Shannon M, Gu Y, Nguyen C;

WPI; 2003-423107/40.

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, associated with decreased or incr MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 1302; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ1, MDZ4, MDZ3, MDZ3, MDZ3, EDZ3, EDZ

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15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3. MDZ4, MDZ7, or MDZ12. e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD23, MD24, MD27, MD212. MD23 is encoded at chromosome [722.1, MD24 is encoded at chromosome [513.2.2, MD27 is encoded at chromosome [611.2 and MD212 is encoded at chromosome [611.2 and MD212 is encoded at chromosome 1611.2 and MD212 is encoded at chromosome or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23, MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic acused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212 genetic locus. The probes are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cytostatic; immunostimulant; gene therapy; vaccine; human;
zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1;
chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                 Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                           Sequence 17 BP; 6 A; 4 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human MDZ3 scanning oligonucleotide SEQ ID 408.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 crdardaacaccadad 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADA99419 standard; DNA; 17
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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD23, MD24, MD212. MD212. MD23 is cancoded at chromosome 7422.1, MD24 is encoded at chromosome 6721.3-22.2, MD27 is encoded at chromosome 6721.3-22.2, MD27 is encoded at chromosome 6721.2 and MD212 is encoded at chromosome cor in manufacturing a medicament for treating or preventing a disorder cor in manufacturing a medicament for treating or preventing a disorder. MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD27, or MD212 genetic locus. The probes are protein in constructing microarrays for measuring gene expression. The proteins are useful at characterize gents for gene therapy or as proteins. The present sequence was used to illustrate the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7g22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15g26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23,
useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
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                                                                                                                  2.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 4.2e+02; tive 0; Mismatches 3; Indels
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                                                                              Sequence 17 BP; 3 A; 8 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human MDZ3 scanning oligonucleotide SEQ ID 404.
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                                                                                                                                                                                                369 ACTITICTIGGACCGCGA 385
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                                                                                                                                                                                                                           1 ACTATCCTGCCCCGCGA 17
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                                                                                                                                                                                                                                                                                                                                       ADA99415 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                             Query Match
Best Local Similarity 82.4°
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                               ADA99415;
                                                                                                                                                                                                                                                                                               RESULT 75
ADA99415
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Score 12.2; DB 1; Length 17; Pred. No. 4.2e+02;

2.9%;

Query Match Best Local Similarity

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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ4, MDZ4, MDZ12, MDZ3 is encoded at chromosome 7422.1, MDZ4 is encoded at chromosome 6721.3-22.2, MDZ7 is encoded at chromosome 6721.2 and MDZ12 is encoded at chromosome 572.2.2, MDZ7 is encoded at chromosome 6721.3 and MDZ12 is encoded at chromosome 672.1 and MDZ12 is encoded at chromosome 1621.2 and MDZ12 is encoded at chromosome 1622 in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased evalopmental disorders. The nucleic acused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross are useful in constructing microarrays for measuring gene expression. The proteins are useful at characteric agents for gene therapy or as proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                               Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                                                                                       Human MDZ3 scanning oligonucleotide SEQ ID 407.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 8; SEQ ID NO 407; 103pp; English.
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                               ADA99418 standard; DNA; 17 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shannon M, Gu Y, Nguyen C;
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                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (AEOM-) AEOMICA INC.
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                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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                                                                               ADA99418;
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     ADA99418
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; Chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
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     Indels
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                                                                                                                                                                                                                                                                                                                                                                                             Human MDZ3 scanning oligonucleotide SEQ ID 405.
  Mismatches
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                                                                                                                                                                                                                                       ADA99416 standard; DNA; 17 BP.
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                                                     365 CCTCACTTTCCTGGACC
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Matches 14; Conservative
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Best Local Similarity
Matches 14; Conserv
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Homo sapiens 3P1281758-A2

3-FEB-2003

20-NOV-2003

ADA99416;

RESULT 79 ADA99416

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Human MDZ4 scanning oligonucleotide SEQ ID 3407.

RESULT 797

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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ12. MDZ3 is encoded at chromosome 7422.1, MDZ4 is encoded at chromosome 6P21.3-22.2, MDZ7 is encoded at chromosome 6P21.3-22.2, MDZ7 is encoded at chromosome 6P21.3-22.2, MDZ7 is encoded at chromosome 16P1.2 and MDZ12 is encoded at chromosome 16P1.2 and MDZ12 is encoded at chromosome 16P1.2 and MDZ12 is encoded at chromosome 16P2.3 and MDZ12 is encoded at chromosome 16P2.3 and MDZ12, or manufacturing a medicament for treating or preventing a disorder or sesociated with decreased or increased expression or activity of MDZ3 MDZ4, MDZ4, MDZ7, or MDZ12. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as proteins. The present sequence was used to illustrate the invention.
                   Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ1; MDZ1; MDZ1; chromosome 7q22.1; chromosome 6p21.3.22.3; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 8; SEQ ID NO 3407; 103pp; English.
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                                                                                                                                                                                                                                                                                                                            32-AUG-2001; 2001US-00922181
                                                                                                                                                                                                                                                                                                                                                                                                                      Shannon M, Gu Y, Nguyen C;
                                                                                                                                                                                                                                                                                                                                                                            (AEOM-) AEOMICA INC.
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                                                                                                                                         Homo sapiens
                                                                                                                                                                                      3P1281758-A2
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ô Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels Seguence 17 BP; 3 A; 5 C; 3 G; 6 T; 0 U; 0 Other;

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Gaps

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HCV DNAzyme substrate sequence #364. ACD57498 standard; RNA; 17 BP (first entry) 23-SEP-2003 ACD57498; RESULT 799 ACD57498 &&&&&&&&X

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Gaps

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Score 12.2; DB 1; Length 17; Pred. No. 4.2e+02; 0; Mismatches 3; Indels

272 GGAGCAGGCGGCACCA 288

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Query Match
Best Local Similarity 82.4%;
Matches 14; Conservative

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ACD58952 standard; RNA; 17 BP

ACD58952;

RESULT 800 ACD58952 ID ACD5895 XX AC ACD5895

Seguence 17 BP; 4 A; 7 C; 6 G; 0 T; 0 U; 0 Other;

Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA sypression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptemer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotopic; cytostatic;

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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed are nucleic acid decory molecules and appearers that bind to HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV cenes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds compounds and/or potential therapies directed against HBV, and compounds methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene expression such as Cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HCV DNAzyme or minus strand DNAzyme sequences disclosed in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                          Lee P;
                                                                                                                                                                                                                                                                                                                                                                                                          Mcswiggen J, Morrissey D, Pavco P,
virucide; antiinflammatory; substrate; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 240; 387pp; English.
                                                                                                                                                                      2001US-00877478
2001US-0296876P
                                                                                                                                                                                                   24-OCT-2001; 2001US-0335059P.
05-DEC-2001; 2001US-0337055P.
                                                                                                                       26-MAR-2002; 2002WO-US009187
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BLATT L.
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Roberts E;
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MCSWIGGEN J.
MORRISSEY D.
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                              Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                            PAVCO P.
LEE P.
DRAPER K.
                                                                                                                                                                                                                                                                                                                                                                            ROBERTS E
                                                           WO200281494-A1
                                                                                                                                                                      08-JUN-2001;
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                                                                                           17-0CT-2002
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Draper K,
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Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; harmarchead ribozyme; DNAzyme; inozyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
             HCV DNAzyme substrate sequence #1090
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Hepatitis C virus.

WO200281494-A1.

17-0CT-2002

26-MAR-2002; 2002WO-US009187

26-MAR-2001; 2001US-00817879. 08-UUN-2001; 2001US-0296876P. 24-0CT-2001; 2001US-0335059P. 05-DEC-2001; 2001US-0337055P.

RIBOZYME PHARM INC.
BLATT L.
MACBCAK D.
MACSHIGGEN J.
MORRISSEY D.
PAVCO P.
LEE P.
LEE P.
LEE P.
LEE P. (RIBO-)
(BLAT/)
(MACE/)
(MCSW/)
(MORR/)
(PAVC/)
(LEEP/)
(DRAP/)

Mcswiggen J, Morrissey D, Pavco P, Macejak D, Roberts E; Blatt L, Me

WPI; 2003-229207/22.

Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus infection.

Claim 1; Page 253; 387pp; English.

The present invention relates to nucleic acid molecules which modulate the synthesis expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymes, DNAzymes, DNAzymes, DNAzymes, and enzymes, zinzymes, amberzymes, and G-cleaver ribozymes, DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes, Also disclosed transcriptase and/or HBV reverse transcriptase primer sequences, as well so oligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV of MBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds complete the expression and/or replication of HCV. The compounds candom and/or useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HCV infection or minus strand DNAzyme sequences disclosed in the present

Sequence 17 BP; 2 A; 5 C; 7 G; 0 T; 3 U; 0 Other;

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Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 70.6%; Pred. No. 4.2e+02; Matches 12; Conservative 2; Mismatches 3; Indels
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Nucleic acid molecule, Hepatitis C virus, HCV; Hepatitis B virus, HBV; RNA stability; RNA expression, RNA synthesis, antisense; enzymatic nucleic acid, hammerhead ribozyme; bNAzyme; inozyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptemen; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                HCV minus strand DNAzyme substrate sequence #1188.
250 CGGGCTCGGCCACGGTG 266
         26-MAR-2001; 2001US-00817879.
08-UUN-2001; 2001US-0087448.
08-UUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0335059P.
05-DEC-2001; 2001US-0337055P.
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BLATT L.
                                                                      ACD63717 standard; RNA; 17
                                                                                                             (first entry)
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MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                                      (RIBO-) RIBOZYME PH.
(BLAT/) BLATT L.
(MACE/) MACBLAK D.
(MCSW.) MCSMIGGEN J.
(MORK/) MORRISSEY D.
(PAVC/) PAVCO P.
(IREP/) LEE P.
(IREP/) LEE P.
(IREP/) RAPERS K.
                                                                                                                                                                                                                                             Hepatitis C virus.
                                                                                                                                                                                                                                                                  WO200281494-A1.
                                                                                                             30-SEP-2003
                                                                                                                                                                                                                                                                                      17-0CT-2002.
                                                                                        ACD63717;
                                                 RESULT 801
ACD63717/c
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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as harmerhead ribozymes, DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV genes and MBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression nucleic acids must be used to modulate of HCV. The compounds that modulate the expression nucleic acids must be settled for restricting and disease states related to HBV and HCV infection, replication and gene and disease states related to HBV and HCV infection, replication and gene Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus Claim 1; Page 296; 387pp; English. infection.

Mcswiggen J, Morrissey D, Pavco P, Lee P;

Macejak D, Roberts E;

Blatt L, N Draper K,

WPI; 2003-229207/22.

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RESULT 803
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense
expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HCV DNAzyme or minus strand DNAzyme sequences disclosed in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for treating cirrhosis, liver failure, oma, or condition associated with hepatitis C virus
                                                                                                                                                                                                                                                                   Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I rection; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                            Gaps
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0
                                                                      Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                  HCV minus strand DNAzyme substrate sequence #2202.
                                                   Sequence 17 BP; 3 A; 7 C; 5 G; 0 T; 2 U; 0 Other;
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                                                                                                                 251 GGGCTCGGCCACGGTGC 267
                                                                                                                                                                                       BP
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08-JUN-2001, 2001US-00877478.
08-JUN-2001, 2001US-0296876P.
24-OCT-2001, 2001US-0335059P.
                                                                                                                                                                                                                                                                                                                                                                                                                             26-MAR-2002; 2002WO-US009187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RIBOZYME PHARM INC. BLATT L.
                                                                                                                                   GGGATCGGTCACCGTGC
                                                                                                                                                                                       ACD65739 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel compound useful for hepatocellular carcinoma,
                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Blatt L, Macejak D,
Draper K, Roberts E;
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MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-229207/22
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DRAPER K.
ROBERTS E.
                                                                                                                                                                                                                                                                                                                                                                 Hepatitis C virus
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                                                                                                                                                                                                                                                                                                                                                                                      WO200281494-A1.
                                                                                                                                                                                                                               30-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Infection
                                    invention
                                                                                                                                                                                                            ACD65739;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MORR/) 1
(PAVC/) 1
(LEEP/) 1
(DRAP/) 1
(ROBE/) 1
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(BLAT/)
(MACE/)
(MCSW/)
                                                                                                                                                                  RESULT 802
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and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, inozymes, zinzymes, amberzymes, and G-claaver ribozymes. Also disclosed transcriptase and/or HBV reverse transcriptase that bind to HBV reverse transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and whethods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HCV brazzie or minus strand DNAzyme sequences disclosed in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid, hammerhead ribozyme; DNAzyme; inozyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; viral replication; virulure; hepatocelular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; 88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.9%; Score 12.2; DB 1; Length 17; 32.4%; Pred. No. 4.2e+02; ve 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HCV minus strand DNAzyme substrate sequence #2024.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mcswiggen J, Morrissey D,
                                                                                                                                                                                                                                                                                                                                                                                                                                               Seguence 17 BP; 2 A; 4 C; 6 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   261 ACGCTGCACCTGGAGCA 277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACD65393 standard; RNA; 17 BP.
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08-UUN-2001; 2001US-0087478.
08-UUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0337055P.
05-DEC-2001; 2001US-0337055P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14; Conservative
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Roberts E;
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MCSWIGGEN J.
MORRISSEY D.
PAVCO P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LEE P.
DRAPER K.
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MACEJAK I
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                                                                                                                                                                                                                                                                                                                                                                                                    invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACD65393;
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(BLAT/)
(MACE/)
(MCSW/)
(MORR/)
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(LEEP/)
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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acid such as hammerhead ribozymes. DNAzymes, inczymes, zinzymes, amberzymes, and G-cleaver ribozymes. DNAzymes are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase primer sequences, as well cranscriptase and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV C genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and compounds compounds and/or potential for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HCV increasion or minus strand DNAzyme sequences disclosed in the present
                                        Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic acid molecule, Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; inozyme; amberzyme; G-dleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; dirhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity 64.7%; Pred. No. 4.2e+02;
11; Conservative 3; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HCV minus strand DNAzyme substrate sequence #1305.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 2 A; 5 C; 5 G; 0 T; 5 U; 0 Other;
                                                                                                                         Claim 1; Page 311; 387pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   129 ATGCTGGCCCGCCTGGC 145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 AUGCUGGCAUUCCUGGC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACD63946 standard; RNA; 17 BP.
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08-JUN-2001; 2001US-0296876P.
08-JUN-2001; 2001US-0335059P.
05-DEC-2001; 2001US-0335059P.
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BLAIT L.
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WPI; 2003-229207/22.
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Best Local Simi:
Matches 11;
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                                                                                     infection.
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(BLAT/)
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The present invention relates to nucleic acid molecules which modulate
the synthesis, expression and/or stability of Hepatitis C virus (HCV) or
Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense
and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes,
inozymes, zinzymes, amberzymes, and G-cleaver ribozymes, Also disclosed
are nucleic acid decoy molecules and aptamers that bind to HBV reverse
transcriptase and/or HBV reverse transcriptase primer sequences, as well
as oligonucleotides that specifically bind the Enhancer I region of HBV
DNA. The nucleic acids may be used to modulate the expression of HBV
genes and HBV viral replication. Also disclosed is a method for screening
compounds and/or potential therapies directed against HBV, and compounds
that modulate the expression and/or periodic for the treatment of degenerative and
disease states related to HBV and HV infection, replication and gene
c expression such as cirrhosas, liver failure, and hepatocellular
carcinoma. The present sequence represents a substrate for one of the HCV
DNAzyme or minus strand DNAzyme sequences disclosed in the present
                                                                                                                                                                                                                                     Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; harmerhead ribozyme; DNAzyme; inozyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptemer; HBV reverse transcriptase; Enhancer I region; viral replication; degemerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
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                                                                                                                                             Mcswiggen J, Morrissey D, Pavco P, Lee
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             y Match 2.9%; Score 12.2; DB 1; Length 17; Local Similarity 70.6%; Pred. No. 4.2e+02; hes 12; Conservative 2; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                   Claim 1; Page 298; 387pp; English.
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                                                                                                                                               Macejak D,
Roberts E;
             MCSWIGGEN J.
MORRISSEY D.
PAVCO P.
LEE P.
DRAPER K.
ROBERTS E.
                                                                                                                                                                                                     WPI; 2003-229207/22
MACEJAK D.
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Draper K,
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                                                     (PAVC/)
(LEBP/)
(DRAP/)
(ROBB/)
MACE/)
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26-MAR-2001; 2001US-00817879.
08-UJN-2001; 2001US-0087478.
08-UJN-2001; 2001US-0296876P.
24-CCT-2001; 2001US-0335059P.
05-DEC-2001; 2001US-0337055P.
                              26-MAR-2002; 2002WO-US009187
                                                                                                                                                                                                                                                                              RIBOZYME PHARM INC.
BLATT L.
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MCSWIGGEN J.
MORRISSEY D.
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DRAPER K.
                                                                                                                                                                                                                                                                                                                                                                                                                                          PAVCO P.
                                                                                                                                                                                                                                                                                                                      BLATT L.
MACEJAK I
                                                                                                                                                                                                                                                                          (RIBO-) | (BLAT/) | (MACE/) | (MCSW/) | (MORK/) | (PAVC/) | (LEEP/) | (DRAP/) | (ROBE/) | (ROBE/
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Mcswiggen J, Morrissey D, Pavco P, Macejak D, Roberts E; Blatt L, M Draper K,

WPI; 2003-229207/22.

Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus infection.

Claim 1; Page 308; 387pp; English.

The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNBzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes, DNBzymes, or nozymes, zinzymes, amberzymes, and G-cleaver ribozymes, Also disclosed transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represente a substrate for one of the HCV invertion.

Sequence 17 BP; 1 A; 6 C; 5 G; 0 T; 5 U; 0 Other;

0; Gaps / Match 2.9%; Score 12.2; DB 1; Length 17; Local Similarity 82.4%; Pred. No. 4.2e+02; les 14; Conservative 0; Mismatches 3; Indels Query Match Best Loca Matches

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ACD62939 standard; RNA; 17 BP ACD62939; RESULT 806 ACD62939
ID ACD6
XX
AC ACD6
XX
DJT 24-8
XX
DZ HCV
XX
NWCl
XX
NWC
XX
NC

(first entry) 24-SEP-2003

HCV minus strand DNAzyme substrate seguence #802.

Nucleic acid molecule, Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme;

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amberzyme; G-cleaver ribozyme; decoy molecule; aptamer;
HBV reverse transcriptaee; Enhancer I region; viral replication;
degenerative; disease state; HBV infection; HCV infection; cirrhosis;
liver failure; hepatocellular carcinoma; hepatotropic; cytostatic;
virucide; antiinflammatory; substrate; ss.
                                                                                                              08-JUN-2001; 2001US-00877478
08-JUN-2001; 2001US-0296876P,
24-OCTI-2001; 2001US-0335059P,
05-DEC-2001; 2001US-0337055P.
                                                                                                        26-MAR-2001; 2001US-00817879
                                                                                         26-MAR-2002; 2002WO-US009187
                                                                                                                                                   RIBOZYME PHARM INC.
BLATT L.
MACBJAK D.
MACSHIGGEN J.
MORRISSEY D.
PAVCO P.
LEE P.
DRAPER K.
ROBERTS E.
                                             Hepatitis C virus
                                                           WO200281494-A1
                                                                          17-0CT-2002.
                                                                                                                                                          (BLAT/)
(MACE/)
(MCSW/)
(MORR/)
(PAVC/)
(LEEP/)
(DRAP/)
(ROBE/)
                                                                                                                                                     (RIBO-)
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ë. Blatt L, Macejak D, Draper K, Roberts E; WPI; 2003-229207/22

Lee

Mcswiggen J, Morrissey D, Pavco P,

Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus infection.

Claim 1; Page 289; 387pp; English.

The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed are uncleic acid decory molecules and G-cleaver ribozymes. Also disclosed are nucleic acid decory molecules and G-cleaver ribozymes. Also disclosed a soligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds charted to the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represente a substrate for one of the HCV DNAzyme or minus strand DNAzyme sequences disclosed in the present invention

Sequence 17 BP; 4 A; 6 C; 7 G; 0 T; 0 U; 0 Other;

ö Query March 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels

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383 CGACGACGCCCAAGA 399 17 1 ceaccacecccaesa ઠે

RESULT 807 ACD64280/c

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ACD64280 standard; RNA; 17
   30-SEP-2003 (first entry)
 ACD64280;
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Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss. ů Mcswiggen J, Morrissey D, Pavco P, Lee HCV minus strand DNAzyme substrate sequence #1471. 26-MAR-2001, 2001US-00817879. 08-UUN-2001, 2001US-00877478. 08-UUN-2001, 2001US-0296876P. 24-0CT-2001, 2001US-0335059P. 05-DEC-2001, 2001US-0337055P. 26-MAR-2002; 2002WO-US009187 RIBOZYME PHARM INC. Roberts E; BLATT L.
MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
PAVCO P.
LEE P.
DRAPER K.
ROBERTS E. Macejak D, WPI; 2003-229207/22 Hepatitis C virus. WO200281494-A1. 17-0CT-2002. Blatt L, N Draper K, (BLAT/) (MACE/) (MCSW/) (MORR/) (PAVC/) (DRAP/) (ROBE/) RIBO-)

Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus infection.

Claim 1; Page 301; 387pp; English.

The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HVV) or Hepatitis B virus (HVV) or The nucleic acid molecules include antisense and enzymatic nucleic acids such as harmer-head ribozymes, DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes, Also disclosed are arenchicase and/or HBV reverse transcriptase primer sequences, as well so oligonucleotides that specifically bind the Enhancer I region of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and methods of the invention and/or replication of HCV. The compounds and disease states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represents a substrate for one of the HCV invention as trand DNAzyme sequences disclosed in the present

T; 1 U; 0 Other; 0 .. ΰ ထ Ä m Sequence 17 BP;

The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes. DNAzymes, inozymes, amberzymes, and G-cleaver ribozymes. Also disclosed are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV genes and HBV viral replication. Also disclosed is a method for screening

Example 1; Page 143; 387pp; English.

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                                                                                                                                                                                                                 Nucleic acid molecule, Hepatitis C virus, HCV; Hepatitis B virus; HBV; RNA stability; RNA expression, RNA synthesis, antisense; enzymatic nucleic acid, hammerhead ribozyme; DNAzyme; inozyme; inozyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antlinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lee
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Length 17;
                       3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mcswiggen J, Morrissey D, Pavco
Query Match
2.9%; Score 12.2; DB 1;
Best Local Similarity 82.4%; Pred. No. 4.2e+02;
Matches 14; Conservative 0; Mismatches 3;
                                                                                                                                                                                               HBV hammerhead ribozyme substrate sequence #361.
                                             140 CCTGGCGGTGGAGGCCG 156
                                                                                                                             BP.
                                                                                                                                                                                                                                                                                                                                                                                                                   26-MAR-2001, 2001US-00817879.
08-UUN-2001, 2001US-00877478.
08-UUN-2001, 2001US-0296876P.
24-0CT-2001, 2001US-03350S9P.
05-DEC-2001, 2001US-03370S5P.
                                                           CCTGGCGGTAGCGGTCG 1
                                                                                                                                                                                                                                                                                                                                                                                                 26-MAR-2002; 2002WO-US009187
                                                                                                                            ACD51048 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RIBOZYME PHARM INC.
                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Macejak D,
Roberts E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
PAVCO P.
LEE P.
DRAPER K.
ROBERTS E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-229207/22.
                                                                                                                                                                                                                                                                                                                          Hepatitis B virus.
                                                                                                                                                                                                                                                                                                                                                  WO200281494-A1.
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                                                                                                                                                                          23-SEP-2003
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Draper K,
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(BLAT/) E
(MACE/) N
(MCSW/) N
(PAVC) I
(PAVC) I
(LEEP/) I
(DRAP/) I
(ROBE/) I
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ACD51048
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10 TGAAACTGCGGGTGACC

Page 414

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compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular earcinoma. The present sequence represents a substrate for one of the HBV riboxyme, inozyme, G-cleaver, zinzyme, DNAzyme or amberzyme sequences disclosed in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cytostatic, virucide, neuroprotective, nootropic, neuroleptic, murine,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Murine oligonucleotide associated with tumour supression, SEQ ID 4145
                                                                                                                                                                                                                                                                                0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                  ch 2.9%; Score 12.2; DB 1; Length 17; I Similarity 58.8%; Pred. No. 4.2e+02; 10; Conservative 4; Mismatches 3; Indels
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                                                                                                                                                                                            Sequence 17 BP; 3 A; 2 C; 6 G; 0 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 515; 738pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tuijnder M;
                                                                                                                                                                                                                                                                                                                      62 GTCTCTGCACTACGAGG 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MOLE-) MOLECULAR ENGINES LAB
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1 GUCUTUGUACUAGGAGG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACC66898 standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-SEP-2001; 2001FR-00011979.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-333167/31.
                                                                                                                                                                                                                                                         Best Local Similarity
Matches 10; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2003025176-A2.
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The present invention relates to murine oligonucleotides (ACC62754-ACC68806), which are associated with tumour suppression, tumour reversion, apoptosis and virus resistance. The oligonucleotides are useful as (1) as probes and primers for detecting, identifying, quantifying and/or amplifying nucleic acid, e.g. as one component of a gene chip, in virco as (anti) sense reagents; and (2) for production of recombinant polypeptides. The oligonucleotides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
                                                                                                                                                                                                                                      Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; ss.
                                                                                                                                                                                                                Murine oligonucleotide associated with tumour supression, SEQ ID 1023
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17 BP; 4 A; 5 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 150; 738pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tuijnder M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   240 GGCTGCTTCCCGGGCTC 256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MOLE-) MOLECULAR ENGINES LAB.
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                           17 recaacreasesreare
                                                                                                               ACC63776 standard; DNA; 17
                                                                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                      Mus musculus.
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                                                                                                                                                 ACC63776;
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                                                                                 RESULT 810
ACC63776/c
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Gaps

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.4-AUG-2003
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Thermus scotoductus; DNA polymerase;
PCR primer; ss.
Phermus scotoductus nucleic acid polymerase PCR primer SEQ ID NO:30.
                                                                  nucleic acid polymerase; enzyme; salt tolerance; thermostability;
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ACC79937 standard; DNA; 17 BP.

RESULT 812

ACC79937;

Synthetic. Thermus scotoductus.

402003066804-A2.

13-SEP-2002; 2002WO-US029102.

14-SEP-2001; 2001US-0322218P. 30-NOV-2001; 2001US-0334489P.

(APPL-) APPLERA CORP. (BOLC/) BOLCHAKOVA E V. (ROZZ/) ROZZELLE J E.

Solchakova EV, Rozzelle JE;

PI; 2003-663590/62

New nucleic acid encoding a Thermus scotoductus strain X-1, ATCC Deposit No. 27978 nucleic acid polymerase, useful for producing nucleic acid polymerases having e.g., improved sequence discrimination or better salt

Example 1; Page 79; 179pp; English

The present invention describes isolated nucleic acids encoding nucleic acid polymerases from Thermus scotoductus. Also described: (1) an isolated nucleic acid (1) encoding a nucleic acid polymerase from Thermus cotoductus strain X-1, ATCC Deposit No. 27978; (2) an isolated DNA polymerase polypeptide from Thermus scotoductus strain X-1, ATCC Deposit No. 27978; (2) an isolated DNA polymerase polypeptide from Thermus scotoductus strain X-1, ATCC Deposit No. 27978; (3) an isolated nucleic acid polymerase; (4) an isolated nucleic acid which encodes a nucleic acid polymerase comprising any of a set of 16 amino acid sequences (82, see ADASOSSOS to ADASO

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Sequence 17 BP; 3 A; 11 C; 1 G; 2 T; 0 U; 0 Other;

0; Gaps Score 12.2; DB 1; Length 17; Pred. No. 4.2e+02; 0; Mismatches 3; Indels Query Match
Best Local Similarity 82.4%;
Matches 14; Conservative (Best Loca Matches

141 CTGGCGGTGGAGGCCGG 157

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Sequencing PCR primer 41 used during construction of B subtilis RB194.

(first entry)

17-0CT-2003

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ABT44053;

BB BB

ABT44053 standard; DNA; 17

17

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The present invention describes a nucleic acid (I) encoding a nucleic acid polymerase or a derivative nucleic acid polymerase with a mutation against decreases 5.3' exonuclease activity or that reduces discrimination against dideoxynucleotide triphosphates. Also described: (1) a vector comprising the nucleic acid (1); (2) a host cell comprising the nucleic acid polymerase or its derivative; (4) a kit acid (1); (3) a nucleic acid polymerase or its derivative; (4) a kit making a container containing the nucleic acid polymerase of (3); (6) synthesising a DNA; (7) thermocyclic amplification of nucleic acid; and (8) primer extending a DNA; The nucleic acid (1) is useful for DNA sequencing or amplification, reverse transcription, RNA amplification or primer extension reactions. The present sequence represents a PCR primer for Thermus oshimai nucleic acid polymerase, which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acid, useful for DNA sequencing or amplification, reverse transcription, RNA amplification or primer extension reactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                  Thermus oshimai; nucleic acid polymerase; enzyme; DNA sequencing; amplification; reverse transcription; RNA amplification; primer extension; PCR primer; ss.
                                                                                                   Thermus oshimai nucleic acid polymerase PCR primer SEQ ID NO:30.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 3 A; 11 C; 1 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 50; 64pp; English.
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                                                                                                                                                                                                                                                                                                                                       22-NOV-2002; 2002WO-US037764.
                                                                                                                                                                                                                                                                                                                                                                         30-NOV-2001; 2001US-0334798P.
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                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                              (APPL-) APPLERA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-505286/47.
                                                                                                                                                                                                                                                                  WO2003048310-A2.
                                                                                                                                                                                                            Thermus oshimai.
Synthetic.
                                                                                                                                                                                                                                                                                                      12-JUN-2003.
                                                                 09-SEP-2003
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ID ABT4

XX ABT4

XX ABT4

DT 17-C

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The invention relates to a novel method which comprises producing a hyaluronic acid via cultivating a Bacillus host cell under conditions suitable for production of the hyaluronic acid and subsequently recovering the hyaluronic acid from the cultivation medium. The most abundant heteropolysaccharides of the body are the glycosaminoglycans, which hyaluronic acid is an example. A number of enzymes are involved in the biosynthesis of hyaluronic acid including hyaluronan synthase, UDP-colucose 6-dehydrogenase, UDP-glucose pyrophosphorylase and UDP-N-colucose of the invention demonstrate ophthalmological, antitrheumatic and dermatcological activities, whilst the method itself may be useful for producing a hyaluronan in a recombinant conthopaedics, rheumatcology or dermatcology and may exhibit further uses within the fields of adhesion, development, cell motility, cancer, orthopaedics, rheumatcology and way exhibit further uses within the fields of adhesion, development, cell motility, cancer, within the fields of adhesion, development esturent sequence is that of the PCR primer of the invention which was used during analysis of the enzymes that play a role in the synthesis of hyaluronic acid
Hyaluronic acid; glycosaminoglycan, hyaluronan synthase; antirheumatic; UDP-glucose 6-dehydrogenase; UDP-glucose pyrophosphorylase; orthopaedic; UDP-N-acetylglucosamine; ophthalmological; dermatological; joint surgery; eye; rheumatology; dermatology; adhesion; development; cell motility; cancer; angiogenesis; wound healing; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Producing a hyaluronic acid (e.g. for use in eye and joint surgery, orthopedics, rheumatology or dermatology) comprises cultivating a Bacillus host cell and recovering the hyaluronic acid from the cultivation medium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sloma A, Behr R, Widner W, Tang M, Sternberg D, Brown S;
                                                                                                                                              Bacillus subtilis subsp. subtilis str. 168.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 11; Page 52; 218pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                            NOVO ) NOVOZYMES BIOTECH INC.
                                                                                                                                                                                                                                                                                                                                                                 21-DEC-2001; 2001US-0342644P.
                                                                                                                                                                                                                                                                                                                20-DEC-2002; 2002WO-US041067.
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                                                                                                                                                                                                                     WO2003054163-A2.
                                                                                                                                                                                                                                                                     03-JUL-2003.
                                                                                                                                                                          Synthetic.
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Gaps . 0 ch 2.9%; Score 12.2; DB 1; Length 17; I Similarity 82.4%; Pred. No. 4.2e+02; 14; Conservative 0; Mismatches 3; Indels Sequence 17 BP; 3 A; 4 C; 5 G; 5 T; 0 U; 0 Other; Query Match Best Local Similarity Best Loca Matches

. 0

84 GCAGTGGACATCACCAC 100 **ccactrcaccrcaacac** δ 요

Tumour suppression/reversion associated nucleotide #4042. ADB43719 standard; DNA; 17 (revised)
(first entry) 18-DEC-2003 04-DEC-2003 ADB43719; RESULT 814 ADB43719/c

BP.

cytostatic; antiviral; neuroprotective; nootropic; neuroleptic; ss; primer; probe; tumour suppression; tumour reversion; apoptosis; virus resistance; transgenic animals; Alzheimer's disease; schizophrenia;

elongation factor 1-alpha; BF-lalpha; pathogen; antibacterial; virucide; fungicide; protozoacide; 88; primer; PCR. Leishmania elongation factor 1-alpha antisense PCR primer SEQ ID NO:20.

Leishmania braziliensis.

WO2003037926-A1

ADC81646 standard; DNA; 17 BP.

ADC81646/ RESULT

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(first entry)

01-JAN-2004

ADC81646;

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The invention relates to the isolation of 6327 nucleotide sequences, cragments of at least 15 consecutive nucleotides of these nucleotides, a sequence having at least 80% identity, after optimal alignment, with the nucleotides, a sequence that hybridizes under stringent conditions with the nucleotides, or the complement, or corresponding RNA, of the confections of the confections, and or used as probes or primers for detecting, cannot seems and antisense sequences, of nucleotides involved in tumour septemental models. The nucleotides involved in tumour corresponding the vectors on a containing them and calls containing the vectors once oncaining them and calls containing the vectors; the encoded polypeptides and antibodies of fural infections or diseases characterized by development of tumours or call degeneration (e.g. Alzheimer's disease or schizophrenia).

Analysis of the expression of the nucleotides can be used for diagnosis and be used to screen for their specific interactive molecules, and polypeptides can be used to screen for their specific interactive molecules.
                                                                                                                                                                                                                                                                                                                                             New nucleic acid encoding human prostate membrane-specific antigen, useful e.g. for treatment of tumors and viral infection, also related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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Pred. No. 4.2e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 5 A; 4 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 504; 771pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         expression of the nucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.9%;
82.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17 Agereereragereare 1
                                                                                                                                                                                                                                                                                                                                                                      useful e.g. for treatment of
polypeptide and antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    400 AGGICTICIACGIGAIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 82.4*
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                          WPI; 2003-441574/41.
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Amson R,

Telerman A,

(MOLE-) MOLECULAR ENGINES LAB

17-SEP-2001; 2001FR-00011981.

17-SEP-2002;

L5-MAY-2003

WO2003040369-A2

Homo sapiens

diagnosis.

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The invention relates to a novel method for the testing of a compound for specific binding to a conserved protein in a pathogen. The method involves comparing binding of the compound to the pathogen. The method of the proteins comprise an insertion/deletion sequence (indel) not present in the other forms of the pathogen and host. The binding to the pathogen form and the absence of or reduced binding of the compound to the host form indicates that the compound is capable of specific binding. A compound of the invention has antibacterial, virucide, fungicide, and proteoracide activity. The method is useful for testing a compound for specific binding to a conserved protein in pathogens (e.g. virus, bacteria, fungi, proteora). The protein is conserved between a pathogen and host (e.g. plant or animal including ammalle e.g. virus, compound is useful as a diagnostic or therapeutic agent specific for the pathogen form, for the preparation of a moiety for specific binding to pathogen form, for the preparation of a moiety for specific binding to pathogen elongation factor lalpha, and for diagnostic and areatment of infectious diseases (e.g. bacterial, viral, fungal and proteozal). The present sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                         Testing of binding specificity of a compound to a conserved protein in a pathogen useful in e.g. treatment of infectious diseases involves comparing binding of the compound to the pathogen and host forms of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Natch 2.9%; Score 12.2; DB 1; Length 17; Local Similarity 82.4%; Pred. No. 4.2e+02; les 14; Conservative 0; Mismatches 3; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene therapy; antibody therapy; modulator of GAPN; GTP-activator for Rab-like GTPase; GAP_N; immunogen; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 1 A; 7 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                  Tcherkassov A, Nandan D;
                                                                                                                                                                                                                                                                                                                                                                                            Example; SEQ ID NO 20; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human GAP_N DNA 17-mer oligo #265
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26 CGAGGCTGGGACGAAG 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       033/c
ADD21033 standard; DNA; 17 BP.
                                                                       01-NOV-2001; 2001CA-02360987.
22-JAN-2002; 2002US-0349319P.
22-JAN-2002; 2002US-0349311P.
05-JUL-2002; 2002US-0393385P.
                                                                                                                                                                          (UYBR-) UNIV BRITISH COLUMBIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-OCT-2002; 2002WO-US032597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-OCT-2001; 2001US-0330323P
                                   01-NOV-2002; 2002WO-CA001689
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17 cdaaddcrdccdaadaa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                     WPI; 2003-482124/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              402003033703-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-APR-2003
                                                                                                                                                                                                                  Reiner NE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADD21033;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                      protein
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Matches
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The invention relates to an isolated human GTP-activator protein for Rab-
like GTPase (GANN) polypeptide (I), a sequence having 65% identity to
(II), a sequence in which at least 95% of deviations from (I) are
conservative substitutions, or a fragment of at least 8 contiguous amino
acids of (I). The polypeptide is useful for identifying a specific
binding partner for itself, by contacting the polypeptide in vivo to a
potential binding partner and determining if the polypeptide binding
partner and determining if the polypeptide binding
partner binds to the polypeptide. (I) and a nucleic acid encoding the
polypeptide (II) are useful for diagnosing or monitoring a disease caused
by altered expression of GAPN, by determining the level of expression of
GAPN in a sample of nucleic acids or proteins that derives from a subject
cuspected to have the disease, alterations from a normal level of
cuspected to have the disease, alterations from a normal level of
cuspected to have the disease, alterations from a normal level of
cuspectated with decreased expression or activity of GAPN, and an
antagonist of (I) is useful for treating or preventing a disorder
associated with increased expression or activity of GAPN, and an
antagonist of (I) is useful for treating or preventing a disorder
associated with increased expression or activity of GAPN, and an
antagonist of (I) is useful for treating or preventing a disorder
cognize GAPN proteins. (II) is useful to drive in vivo expression of
GAPN proteins, and as hybridization probes to detect, characterize and
cuantify GAPN nucleic acids in and isolate GAPN nucleic acids from both
genomic and transcript-derived nucleic acids samples. This sequence.

The presents a 17-mer oligonucleotide spanning the GAP. NON sequence.
                                                                                                                                                     Novel human GTP-activator protein for Rab-like GTPase and polynucleotide encoding the protein, useful for diagnosing, treating or preventing disorders associated with increased expression or activity of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3, Indels : 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.9%; Score 12.2; DB 1; Length 17; 12.4%; Pred. No. 4.2e+02; ve 0; Mismatches 3; Indels :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene therapy; antibody therapy; modulator of GAPN; GTP-activator for Rab-like GTPase; GAP_N; immunogen; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17 BP; 2 A; 7 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                       Example 2; SEQ ID NO 289; 149pp; English.
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                     (AMSH ) AMERSHAM BIOSCIENCES SV CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human GAP_N DNA 17-mer oligo #115.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     254 CTCGGCCACGGTGCACC 270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADD20883 standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17 cacadecacacidatecrec 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 2.9%;
Best Local Similarity 82.4%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                            WPI; 2003-403224/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2003033703-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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                                                                   Zhang J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    817
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ADD20883
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Gaps ·,

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Zhang J;
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WPI; 2003-403224/38.

Novel human GTP-activator protein for Rab-like GTPase and polynucleotide encoding the protein, useful for diagnosing, treating or preventing disorders associated with increased expression or activity of the protein.

Example 2; SEQ ID NO 139; 149pp; English

The invention relates to an isolated human GTP-activator protein for Rablike GTPase (GAPN) polypeptide (I), a sequence having 65% identity to
(I), a sequence in which at least 95% of deviations from (I) are
conservative substitutions, or a fragment of at least 8 contiguous amino
acids of (I). The polypeptide is useful for identifying a specific
binding partner for itself, by contacting the polypeptide in vivo to a
potential binding partner and determining if the polypeptide binding
partner binds to the polypeptide (I) and a nucleic acid encoding the
polypeptide (II) are useful for diagnosing or monitoring a disease caused
by altered expression of GAPN, by determining the level of expression of
GAPN in a sample of nucleic acids or proteins that derives from a subject
cuspected to have the disease, alterations from a normal level of
expression providing diagnostic and/or monitoring information. (I), (II)
cor agonist of (I) is useful for treating or preventing a disorder
associated with increased expression or activity of GAPN, and an
antegonist of (I) is useful for treating or preventing a disorder
associated with increased expression or activity of GAPN, and an
antegonist of (I) is useful for treating or preventing a disorder
cor agonist of (I) is useful for treating or preventing a disorder
associated with increased expression or activity of GAPN, and an
antegonist of (I) is useful for treating or preventing a disorder
cor agonist of (I) is useful for treating or preventing a disorder
associated with increased expression or activity of GAPN proteins.
(I) is useful as immunogen to raise antibodies that specifically
recognize GAPN proteins. (II) is useful to drive in vivo expression of
genomic and transcript-derived mucleic acids samples. This sequence

Sequence 17 BP; 3 A; 6 C; 6 G; 2 T; 0 U; 0 Other;

Gaps ÷ 2.9%; Score 12.2; DB 1; Length 17; 82.4%; Pred. No. 4.2e+02; tive 0; Mismatches 3; Indels Query Match
Best Local Similarity 82.44
Matches 14; Conservative

ADD20884 standard; DNA; 17 ADD20884; RESULT 818 ADD20884

(first entry) 15-JAN-2004

fuman GAP_N DNA 17-mer oligo #116

gene therapy; antibody therapy; modulator of GAPN; GTP-activator for Rab-like GTPase; GAP_N; immunogen; ss.

Homo sapiens.

WO2003033703-A2.

24-APR-2003

11-OCT-2002; 2002WO-US032597

.5-OCT-2001; 2001US-0330323P

(AMSH) AMERSHAM BIOSCIENCES SV CORP.

Zhang J;

WPI; 2003-403224/38.

Novel human GTP-activator protein for Rab-like GTPase and polynucleotide encoding the protein, useful for diagnosing, treating or preventing disorders associated with increased expression or activity of the

Example 2, SEQ ID NO 140; 149pp; English.

The invention relates to an isolated human GTP-activator protein for Rablike GTPase (GAPN) polypeptide (I), a sequence having 65% identity to (I), a sequence in which at least 95% of deviations from (I) are conservative substitutions, or a fragment of at least 8 contiguous amino acids of (I). The polypeptide is useful for identifying a specific binding partner for itself, by contacting the polypeptide in vivo to a potential binding partner and determining if the polypeptide binding partner and determining the level of expression of GAPN, by altered expression of GAPN, by determining the level of expression of ample of nucled acids or proteins that derives from a subject suspected to have the disease, alterations from an ormal level of expression or agonist of (I) is useful for treating or preventing a disorder associated with increased expression or activity of GAPN, and an antagonist of (I) is useful for treating or preventing a disorder associated with increased expression or activity of GAPN and an antagonist of (I) is useful to raise antibodies that specifically recognize GAPN proteins. (II) is useful to drive in vivo expression of GAPN proteins, and as hybridization probes to detect, characterize and quantify GAPN nucleic acids in and isolate GAPN nucleic acids from both genomic and transcript-derived mucleic acids samples. This sequence.

Sequence 17 BP; 3 A; 7 C; 5 G; 2 T; 0 U; 0 Other;

Gaps ô 2.9%; Score 12.2; DB 1; Length 17; 32.4%; Pred. No. 4.2e+02; [ve 0; Mismatches 3; Indels Query Match 2.9%; Best Local Similarity 82.4%; Matches 14; Conservative

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ADD21031 standard; DNA; 17 BP. RESULT 819 ADD21031/c

(first entry) 15-JAN-2004 ADD21031;

Human GAP_N DNA 17-mer oligo #263

gene therapy, antibody therapy; modulator of GAPN; GTP-activator for Rab-like GTPase; GAP_N; immunogen; ss.

Homo sapiens

WO2003033703-A2.

24-APR-2003,

11-OCT-2002; 2002WO-US032597

15-OCT-2001; 2001US-0330323P.

(AMSH) AMERSHAM BIOSCIENCES SV CORP.

Zhang J;

WPI; 2003-403224/38

Novel human GTP-activator protein for Rab-like GTPase and polynucleotide encoding the protein, useful for diagnosing, treating or preventing disorders associated with increased expression or activity of the protein.

Example 2; SEQ ID NO 287; 149pp; English.

The invention relates to an isolated human GTP-activator protein for Rablike GTPase (GAPN) polypeptide (I), a sequence having 65% identity to (I), a sequence in which at least 50° f deviations from (I) are conservative substitutions, or a fragment of at least 8 contiguous amino acids of (I). The polypeptide is useful for identifying a specific binding partner for itself, by contacting the polypeptide in vivo to a perential binding partner and determining if the polypeptide binding partner and determining if the polypeptide binding partner by the polypeptide (II) are useful for diagnosing or monitoring a disease of SAPN in a sample of mucleic acids or proteins that derives from a subject subspected to have the disease, alterations from a normal level of expression providing diagnostic and/or monitoring a disorder associated with decreased expression or activity of GAPN, and an associated with decreased expression or activity of GAPN, and an anagonist of (I) is useful for treating or preventing a disorder associated with increased expression or activity of GAPN, and an anagonist of (I) is useful for treating or preventing a disorder associated with increased expression or activity of GAPN, and an anagonist of (I) is useful for treating or preventing a disorder associated with increased expression or activity of GAPN, and an anagonist of (I) is useful for treating or preventing a disorder associated with increased expression or activity of GAPN and an anagonist of (I) is useful for treating or preventing a disorder associated with increased expression or activity of GAPN proteins. (II) is useful to drive in vivo expression of quantify GAPN nucleic acids in and isolate GAPN nucleic acids in and isolated. This sequence represents a 17-mer oligonucleotide spanning the GAP N DNA sequence.

Sequence 17 BP; 2 A; 7 C; 6 G; 2 T; 0 U; 0 Other;

ô 0; Gaps Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels

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RESULT 820

Human GAP_N DNA 17-mer oligo #117. ADD20885 standard; DNA; 17 BP (first entry) 15-JAN-2004 ADD20885; ADD20885

gene therapy; antibody therapy; modulator of GAPN; GTP-activator for Rab-like GTPase; GAP_N; immunogen; ss.

Homo sapiens.

WO2003033703-A2.

24-APR-2003,

11-OCT-2002; 2002WO-US032597.

15-OCT-2001; 2001US-0330323P.

(AMSH) AMERSHAM BIOSCIENCES SV CORP

Zhang J;

WPI; 2003-403224/38.

Novel human GTP-activator protein for Rab-like GTPase and polynucleotide

encoding the protein, useful for diagnosing, treating or preventing disorders associated with increased expression or activity of the protein.

Example 2; SEQ ID NO 141; 149pp; English.

The invention relates to an isolated human GTP-activator protein for Rab
(1), a sequence in which at least 95¢ of deviations from (1) are

(1), a sequence in which at least 95¢ of deviations from (1) are

conservative substitutions, or a fragment of at least 8 contiguous amino

acids of (1). The polypeptide is useful for identifying a specific

binding partner for itself, by contacting the polypeptide binding

partner binds to the polypeptide. (1) and a nucleic acid encoding the

potential binding partner and determining if the polypeptide binding

partner binds to the polypeptide. (1) and a nucleic acid encoding the

polypeptide (II) are useful for diagnosing or monitoring a disease caused

by altered expression of GAPN, by determining the level of expression of

CAPN in a sample of nucleic acids or proteins that derives from a subject

caspociated with decreased and/or monitoring information. (1), (II)

caspociated with decreased expression or activity of GAPN, and an

antagonist of (1) is useful for treating or preventing a disorder

caspociated with increased expression or activity of GAPN, and an

antagonist of (1) is useful for treating or preventing a disorder

caspociated with increased expression or activity of GAPN, and an

antagonist of (1) is useful for treating or preventing a disorder

caspociated with increased expression or activity of GAPN, and an

antagonist of (1) is useful for treating or preventing a disorder

caspociated with increased expression or activity of GAPN, and an

cantagonist of (1) is useful for treating or preventing a disorder

caspociated with increased expression or activity of GAPN, and an

caspociated with increased expression or activity of GAPN, and an

cantagonist of (1) is useful for treating or preventing a disorder

caspociated with increased expression or activity of GAPN, and an

cappeted to a second and proper to raise antibodies that specifically

crecopal and proper active or a second and an expecification probes to detect, characterize and

cappeted to activ

Seguence 17 BP; 3 A; 8 C; 4 G; 2 T; 0 U; 0 Other;

Gaps ö Query Match 2.9%; Score 12.2; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 4.2e+02; Matches 14; Conservative 0; Mismatches 3; Indels

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AAQ22266 standard; DNA; 18 RESULT 821

20-JUL-1992 (first entry)

AAQ22266;

Methylphosphonate oligomer #0059 complementary to HSV-1 polyA signal.

Herpes Simplex Virus; type 1; beta-gene; UL5; DNA dependent ATPase; ss.

Synthetic.

WO9203051-A

05-MAR-1992.

90US-00568501, 15-AUG-1990;

90US-00568501. (GENT-) GENTA INC. 15-AUG-1990; X2X5X8X8X8X8X8X8X8X8X8X8X8X8XX

Roizman B, Maxwell KW;

WPI; 1992-096516/12.

New oligomers complementary to viral genome(s) or mRNA transcripts areanti-sense agents which interfere with viral replication of e.g. Herpes simplex virus, Epstein-Barr virus etc.

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Example 2; Page 20; 33pp; English.
                                   also AAQ22247-Q22283
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This oligomer contains methylphosphonate linkages except for the first 5' linkage which is a phosphate diester bond. The oligomer is complementary to the area around the polyA signal of the HSV-1 ULS gene. ULS is one of the essential beta-genes and the protein it encodes forms a complex with two other proteins which functions as a primase and helicase. The protein specified by ULS has also been shown to act as a DNA dependent ATPase. The oligomer can interfere with expression and function of the gene. See

Sequence 18 BP; 2 A; 9 C; 4 G; 3 T; 0 U; 0 Other;

ö Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels

327 GCGGCGGACGACCAGGG 343 18 GCGACGGCGATCAGGG 2 g ð

AAQ41689 standard; DNA; 18 AAQ41689; 822 AAQ41689 RESULT

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(revised)
(first entry) 25-MAR-2003 24-AUG-1993

for Class I HLA DNA allele B region C. Probe RAP14 Amplification; allelic variants; A; B; C; alleles; exon; diagnosis; tissue typing; forensic testing; susceptibility; PCR; ss.

Synthetic.

28-OCT-1992; 12-MAY-1993

92EP-00118396,

HOFF) HOFFMANN LA ROCHE & CO AG

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Bugawan T, Erlich HA;

WPI; 1993-153998/19.

Rapid HLA class I typing of sample nucleic acid - by amplifying second or third exon sequences then hybridising with set of specific probes, useful e.g. in tissue typing and forensic tests.

Disclosure; Page 7; 23pp; English.

The HLA class I DNA type of nucleic acid in a sample may be determined by amplifying any DNA contg. a class I HLA allele second and/or third exon, hybridising the PCR prod. with probes which only hybridise to exactly complementary sequences and detecting the pattern of hybridisation given, which is indicative of the Class I HLA allele of the sample. The A, B and opt. C alleles are amplified by PCR using pairs of nucleotide primars. Specific primars for exon 2 are D8308 and D8309 and for the third exon are D8311 and D837. A panel of sequence specific oligonucleotide probes (SSO8) is used to detect the HLA A and B allelic variants not distinguishable by serological, cellular or biochemical methods. The region identifies the polymorphic codons of the second exon of Class I HLA A or B alleles to which the probe hybridises. Region A includes codons codons 30 f exon 2 of A alleles. Region D includes codons 62 and 63 of exon 2 of A alleles. Region D includes codons 73

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-77 of exon 2 of A alleles. Specific applications include tissue typing, identification of individuals (e.g. in forensic tests) and detecting susceptibility to disease. See also AAQ41656-94. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                       Gaps
                                                                                                                                                                       .
0
                                                                                                                                2.9%; Score 12.2; DB 1; Length 18; 82.4%; Pred. No. 4.7e+02; tive 0; Mismatches 3; Indels
                                                                                                5 G; 1 T; 0 U; 0 Other
                                                                                                                                                                                                             174 TACGAGTCCAAGGCACA 190
                                                                                                                                                                                                                                                 1 racaaeeeccaeeeaca 17
                                                                                                  Sequence 18 BP; 6 A; 6 C;
                                                                                                                                  Query Match 2.9
Best Local Similarity 82.4
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                            RESULT 823
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BP.

AAQ53969 standard; DNA; 18

AAQ53969 ID AAQ5

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Gaps

(first entry)

03-AUG-1995

AAQ53969;

Human, OTC; identification; mutation; amplify; PCR; diagnosis; fluoresence-label; primer; electrophores; genetic disease; single stranded conformation polymorphism; SSCP; detection; ss. Human OTC gene sense primer, binds to bases 21-39. JP05317048-A. 30-SEP-1992; 03-DEC-1993 Synthetic

91JP-00280835. (SHIO) SHIONOGI & CO LTD. (MATS/) MATSUDA I. 30-SEP-1991;

WPI; 1994-011017/02.

Gene mutation identification for genetic disease diagnosis - includes specific gene or fragment amplification by polymerase chain reaction using fluorescence-labelled primer and electrophoresising.

Disclosure, Page 12; 14pp; Japanese

The sequences given in AAQS3956-78 are primers which were used in the method of the invention to detect mutations in the human OTC gene. The gene is amplified by PCR using a fluoresence-labelled primer and the amplified gene or fragment is electrophoresed by single stranded conformation polymorphism (SSCP) and detecting the mutated gene via the primer. This method can be used to detect the presence of mutation in a perme with a preciseion equal to or higher than that of RI-labelling methods. This method may be used in the diagnosis of genetic disease

Sequence 18 BP; 6 A; 4 C; 6 G; 2 T; 0 U; 0 Other;

Gaps ö Query Match
2.9%; Score 12.2; DB 1; Length 18;
Best Local Similarity 82.4%; Pred. No. 4.7e+02;
Matches 14; Conservative 0; Mismatches 3; Indels

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RESULT 824 AAT05082

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AAT05082 standard; DNA; 18

WO9720197-A2

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A novel method has been developed for identifying an unknown allele of a polyallelic gene. The method involves: (a) contacting the unknown allele polyallelic gene. The method involves: (a) contacting the unknown allele converted by a panel of probes, each of which recognises a sequence motif that is present in some alleles of the polyallelic gene but not in others; (b) coherving which probes recognise the unknown allele so as to obtain a cingerprint of the unknown allele; and (c) comparing the fingerprint with fingerprints of the unknown allele; and (c) comparing the fingerprint with cingerprints of known alleles. The present sequence represents a cingerprint of the unknown allele. The method where the polyallelic specifically claimed probe for use in the method where the polyallelic concepts as human enthocyper, and genes, specifically the HLA class I and complement factor genes I and CC. If genes the T cell receptor genes in mammals, TAP, LMP, ras, conclassical HLA class I genes, human complement factor genes C4 and C2, conclassical that complex, and genes located in mitochondrial DNA, bacterial corrections and viral DNA. The method is particularly useful for matching conception to tissue or organ transplantations. The method can also be used in paternity testing, in forensic medicine, as a follow up technique cused in paternity testing, in forensic medicine, as a follow up technique adoptive immunotherapy, and in identification of alleles of the concept in intended recurring motif conception of polyallelic genes using a limited number of selected recurring motif
                                                                                                                                                                                                                                                                                                                                                    Identifying unknown allele(s) of a poly:allelic gene using panel of probes each recognising a sequence motif present in some allele(s) useful for donor matching in tissue transplantation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 2 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                  (NOLA-) NOLAN BONE MARROW TRUST ANTHONY.
                                                                                                                                                                                                                                                       Madrigal A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 5; Page 19; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     298 AGGACCTGAGCCCCGGG 314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAX75558 standard; RNA; 18 BP.
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                                                                                                96WO-GB002959
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                                                                                                                                                                                                                                                    Arguello R, Avakian H,
                                                                                                                                                                                                                                                                                                WPI; 1997-310717/28.
                                                                                                29-NOV-1996;
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                                              05-JUN-1997.
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AAX75558
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A PCR primer pair (AAT05082-83) correspond to a sense sequence in exon 2 of HLA-A1 antigen and an antisense sequence in exon 3, respectively. The parimers were used in PCR and RT-PCR with tumour rejection antigen precursor MAGE gene-based primers to detect MAGE gene expression in tumours and normal tissues. (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Determn. of cancerous condition(s) - using a nucleic acid as a primer to determine expression of a MAGE tumour rejection antigen precursor.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           De Plaen E, Boon-Falleur T, Lethe B, Szikora J, De Smet C;
Chomez P, Gaugler B, Van Den Eynde B, Brasseur F, Patard J;
Weynants P, Marchand M, Van Der Bruggen P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels
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                                                                                                                                                                                                                              MAGE; tumour rejection antigen; cancer; diagnosis; polymerase chain reaction; PCR; primer; HLA-Al; ss
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                                                                                                                                                                            ILA-A1 PCR primer (sense, exon 2).
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94US-00209172.
94US-00299849.
94US-00346774.
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                                                                                                (revised)
(first entry)
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10-MAR-1994;
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30-NOV-1994;
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Weynants P,
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26-FEB-1996
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                                                                                                                                                                                                                                                                                                           Synthetic.
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                                                                                                                                                                                                                                                                                                                 Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour anglogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-1ike tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; 88.
                                     Gaps
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Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                      Mouse flt-1 VEGF receptor hairpin ribozyme substrate #17.
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Human leukocyte antigen; HLA; probe; tissue transplantation; MHC gene; major histocompatibility complex; paternity test; forenaic medicine; haematological malignancy; inherited disorder; adoptive immunotherapy; identification; ss.

Homo sapiens

Synthetic

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Folkerts O,
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                            WPI; 1997-202224/18.
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08-DEC-1998
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Young SA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAV60768;
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                                                                                                                                                                                                                                                                                                                        The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGP). A patient (preferably human) having a condition associated with the level of the fims-like tyzosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour anglogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                              Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Maize; corn; Zea mays; delta-9 desaturase; GBSS; target; substrate; granule bound starch synthase; hammerhead ribozyme; hairpin ribozyme; modulation; gene expression; transgenic plant; cleavage; canola plant; caffeine synthesis; coffee plant; nicotine production; tobacco; fruit ripening; flower pigmentation; lignin production; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Skokut TA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Granule bound starch synthase hairpin substrate SEQ ID NO:635.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 70.6%; Pred. No. 4.7e+02; Matches 12; Conservative 2; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Merlo PAO,
                                                                                                                                                                     Escopedo J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seguence 18 BP; 3 A; 6 C; 6 G; 0 T; 3 U; 0 Other;
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                                                                                                                                                                     Stinchcomb D,
                                                                                                                                                                                                                                                                                                Claim 4; Page 185; 218pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              238 GAGGCTGCTTCCCGGGC 254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 GAGACUGCUCCACGGGC 18
                                                                          95US-0005974P.
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                                            96WO-US017480.
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                                                                                                                         PHARM INC.
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                                                                                                                                                                     Pavco P, Mcswiggen J,
                                                                                                                                                                                                     WPI; 1997-259017/23.
                                                                                                                     (RIBO-) RIBOZYME PHA
(CHIR ) CHIRON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-JUL-1995;
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                                                                          26-OCT-1995;
11-JAN-1996;
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                                            25-OCT-1996;
                01-MAY-1997.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX62760;
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                                                                                                                                                                                                                                                                                                                                               with RNA cleaving activity, which modulates the expression of a plant gene. Also described is a gene comprising a cDNA sequence encoding maize Delta-9 desaturase. (I) can be used to modulate expression of a gene, preferably Delta-9 desaturase or a granule bound starch synthase (GBSS) gene, in a plant (preferably a maize or canola plant). (I) can be used to modulate expression of a gene, in a plant (preferably a maize or canola plant). (I) can be used to tobacco plant, fruit ripening processes in an apple, tomato, pear, plum or peach plant, flower pigmentation in a rose, petunia, chrysanthemum or marigold plant or lignin production in a tobacco, aspen, poplar or pine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Non-M, non-O HIV-1 strain YBF30 - useful for diagnosis and immunisation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HIV-1 strain YBF30; antibody; oligonucleotide; diagnosis; immunisation; infection; typing; gag; PCR primer; amplification; ss.
                                                                                                                                  Ribozyme which modulates plant gene expression - preferably modulates expression of DELTA-9 desaturase or granule bound starch synthase in maize or canola.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Y Match 2.9%; Score 12.2; DB 1; Length 18; Local Similarity 70.6%; Pred. No. 4.7e+02; he 12; Conservative 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 18 BP; 2 A; 6 C; 7 G; 0 T; 3 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HIV-1 strain YBF30 gag gene primer GAG Y S1.1.
                                                                                                                                                                                                                                                                                        Claim 42; Page 84; 155pp; English.
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Human immunodeficiency virus 1.
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Merlo
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(first entry)
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peptides, antibodies and oligonucleotides derived from it (see AAV60752-
V60798 and AAW68473-W68482) are used for diagnosis of or immunisation
against non-M, non-O HIV-1 infections. The oligonucleotides, peptides and
antibodies can also be used for typing HIV strains. (Updated on 25-WAR-
2003 to correct PI field.)
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Sequence 18 BP; 6 A; 5 C; 3 G; 4 T; 0 U; 0 Other;

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Gaps
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Match 2.9%; Score 12.2; DB 1; Length 18; Local Similarity 82.4%; Pred. No. 4.7e+02; les 14; Conservative 0; Mismatches 3; Indels
     Query Match
                                        Matches
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71 55 CAGAGGAGTCTCTGCAC

CAGAGACTCTCTGTAC 18 셤

AAV34526 standard; DNA; 18 BP 829 RESULT 82 AAV34526/

AAV34526;

Chemokine receptor CXCR4 amplifying RT-PCR primer 2.

(first entry)

20-AUG-1998

Chemokine receptor; gp120; fusion protein; HIV; screening; AIDS; CD4 binding site; RT-PCR primer; ss.

Homo sapiens Synthetic

409815569-A1

16-APR-1998

97WO-US018397. 08-OCT-1997; 96US-0027931P. 19-OCT-1996; DAND) DANA FARBER CANCER INST INC.

(LEUK-) LEUKOSITE INC. (CHIL-) CHILDRENS MEDICAL CENT

WPI; 1998-240778/21.

Gerard C;

Gerard N,

Wu L,

Newman W,

Sodroski J,

Derivatives of gp120 containing modified chemokine receptor binding site - and complexes with soluble CD40, for inhibiting infectivity of human immune deficiency virus and to screen for inhibitors.

Example; Page 53; 92pp; English

This primer is used for the RT-PCR amplification of a chemokine receptor CXCR4. The invention provides gpl20 derivative having a conformational, discontinuous chemokine receptor binding site defined by amino acids residues present in the gpl20 constant regions C2, C3 and C4, and the variable region V3, and its conformation is similar to that of the creeptor binding site of wild-type gpl20 complexed to CD4. Exposure of the chemokine receptor binding site is increased by having at least part of a variable or constant region of wild-type gpl20 removed. A stabilised complex of gpl20 CD4 binding site with a soluble CD4 molecule is used to inhibit infectivity of human immune deficiency virus (HIV). Labelled gpl20 derivatives are used to screen for inhibitors of HIV infection from increased levels of the chemokine receptors (at the chemokine receptor are used as models for diagnosing susceptibility complexed in or undistant of an increased levels). Transgenic animals expressing CD4 and chemokine receptor are used as models for studying development of AIDS or effect/safety of therapeutic agents

Sequence 18 BP; 4 A; 4 C; 7 G; 3 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAV46054 and AAV46200-V46264 are primers used in isolating human histocompatibility locus antigen (HLA-A) Class I alleles which are used in a novel method of HLA-Class I typing. The method involves combining a group-specific untranslated region primer pair with a target DNA to allow primer-based amplification of the DNA, and determining whether a nucleic acid product is produced by the amplification. The ability of the primer pair to produce a product is associated with a particular HLA group type. The methods can be used for typing the 3 classical HLA Class I genes (comprising the loci HLA-A, HLA-B, and HLA-C) in e.g. donors and hosts for tissue transplantation. The initial group specific amplification allows a PCR based separation of haplotypes in 95% of patient samples. The subsequent sequencing can provide for high-resolution typing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIA Class I typing - by primer-based amplification of target DNA using group-specific untranslated region primer pair.
                                                                                                                                                                                                                                                                                                      Histocompatibility locus antigen; HLA-A class I; human; class typing; donor; host; tissue transplantation; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gape
                                 Gaps
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 Length 18;
                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 1 A; 7 C; 6 G; 4 T; 0 U; 0 Other;
2.9%; Score 12.2; DB 1;
82.4%; Pred. No. 4.7e+02;
tive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 8; Page 138; 185pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV39316 standard; cDNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            230
                                                                343 GCCGGCTGCTCTACAGC 359
                                                                                                                                                                             BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                96US-00766189.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (VISI-) VISIBLE GENETICS INC.
                                                                                              17 geresereceraciae 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV46248 standard; DNA; 18
                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Leushner J;
                                 14; Conservative
                                                                                                                                                                                                                                                                        Human HLA-A primer #152.
   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1998-348544/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Blasczyk RH,
                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                   WO9826091-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-DEC-1996;
                                                                                                                                                                                                                                          16-0CT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                 18-JUN-1998
                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                          AAV46248;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 831
AAV39316/c
ID AAV393
XX
                                                                                                                                               RESULT 830
                                   Matches
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Wed Apr 21 12:58:21 2004
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The present sequence represents a PCR primer for use in a method of the invention for determining the genetic predisposition to cancer in an invention for determining the genetic predisposition to cancer in an invention for determining the genetic predisposition to cancer in an invention of processent in tumnours that display allelic imbalance at Ip32, the chromosomal band identified as one of four minimal regions of chromosomal deletion in breast carcinomas. hRAD54 is useful for production of proteins, inter alia, that have been identified as novel haxD54 by homology between the amino acid sequence given in AAW62186 and known amino acid sequences such as yeast RAD54. hRAD54 proteins are used in the treatment of cancer, including Xeroderma Pigmentosum and Bloom syndrome, Werner's syndromes and breast cancer. hRAD54 polynucleotides are also useful for detecting complementary nucleotides for use as a diagnostic agent, especially useful for diagnosis of disease or susceptibility to diseases. hRAD54 polynucleotide, proteins, agonists and antagonists which Human hRAD54 DNA and polypeptide - and agonists, antibodies, antagonists, Human; RAD54; hRAD54; cancer; xeroderma pigmentosum; Bloom syndrome; Werner's syndrome; ATR-X; diagnosis; detection; SNF2 superfamily; X-linked mental retardation with alpha-thalassemia syndrome; tumour; gene therapy; PCR primer; ss. Human RAD54 mutation detecting PCR primer SEQ ID NO:24. Robbins DJ; Claim 18; Page 39; 64pp; English Rasio D, (SMIK) SMITHKLINE BEECHAM CORP. (UYJE-) UNIV JEFFERSON THOMAS. 97EP-00308998. 96US-0030676P. (first entry) Croce CM, Fishel RA, WPI; 1998-274189/25 .3-NOV-1996; Homo sapiens 10-NOV-1997; 16-SEP-1998 EP844305-A2 Synthetic AAV39316;

Sequence 18 BP; 3 A; 5 C; 5 G; 5 T; 0 U; 0 Other;

ö Score 12.2; DB 1; Length 18; Pred. No. 4.7e+02; 3; Indels 0; Mismatches Query Match
Best Local Similarity 82.4%;
Matches 14; Conservative

286 CCAAGCTGGTGAAGGAC 302

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18 ccágcchéchcaaca

Angiogenin antisense oligonucleotide JF12S.

Angiogenin; antisense; inhibitor; cancer; metastasis; angiogenesis;

New oligo:nucleotide(s) that inhibit expression of angiogenin - for treatment of tumours and metastases, or other conditions involving abnormal angiogenesis. Location/Qualifiers
1.18 /*ta.9 a /*toce "phosphorothioate linkages" //note= "phosphorothioate linkages" Claim 10; Page 38; 71pp; English. 98WO-US005651. (HARD) HARVARD COLLEGE WPI; 1998-531944/45. Olson KA; Key modified_base 20-MAR-1998; WO9842722-A1 21-MAR-1997; 01-OCT-1998 Synthetic Pett JW,

Antisense phosphorothioate oligonucleotide JF10S encompasses the AUG initiation codon of the human angiogenin gene (see AAV60918). JF2S, and other claimed antisense oligonucleotides (see AAV60911) with base sequences complementary to a target region of the angiogenin gene, are able to inhibit expression of angiogenin. They are used in claimed methods to decrease production of angiogenin, particularly to reduce the size of tumours associated with angiogenesis, to inhibit metastases, establishment of tumour cells or growth of tumours and, when labelled, to detect angiogenein for diagnosis of conditions associated with abnormal angiogenesis. They can also be used to treat a wide range of non-cancer conditions that involve angiogenesis, e.g. age-related macular degeneration, diabetic retinopathy, bacterial or fungal ulcers, many others listed

Sequence 18 BP; 4 A; 9 C; 3 G; 2 T; 0 U; 0 Other;

Gaps .. 0 Length 18; Score 12.2; DB 1; Length 18 Pred. No. 4.7e+02; 0; Mismatches 3; Indels Query Match 2.9%; Best Local Similarity 82.4%; Matches 14; Conservative

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292 TGGTGAAGGACCTGAGC 308 17 regrearedeccreeec 셤 à

AAZ11707 standard; RNA; 18 BP. AAZ11707;

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Gaps

32-NOV-1999 (first entry)

bases 330-347. Hepatitis C virus antisense DNA 25 - binds to HCV genome

Antisense; oligonucleotide; hepatitis C virus; antiviral detection; diagnosis; treatment; translation inhibition; replication inhibition; ss.

Synthetic. Hepatitis C virus.

WO9929350-A1

therapy, diagnosis, ss.

Nozaki C;

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Hepatitis C virus antisense DNA 34 - binds to HCV genome bases 331-348.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense; oligonucleotide; hepatitis C virus; antiviral therapy; detection; diagnosis; treatment; translation inhibition; replication inhibition; ss.
                                                                                                                                                                                                                                                                                                 Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                 261 ACGGTGCACCTGGAGCA 277
                                                                                                                                                                                                                                                                                                                                                                                      AAZ11716 standard; RNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98WO-US026040.
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                                                                                                                                                                                                                                                                                                                                                Accerecaceareages 2
                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                   (ISIS-) ISIS PHARM INC
                                                                                 IPI; 1999-493767/41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
Hepatitis C virus.
                                                                                                                                                                                                                                                                      other agents
                                                                   Anderson KP,
                                                                                                                                                                                                                                                                                                                                                                                                                    02-NOV-1999
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                      08-DEC-1998;
                                    10-DEC-1997;
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       17-JUN-1999
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                                                                                                        infections
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Human; PRO; EST; expressed sequence tag; PCR primer; hybridisation; probe; blood coagulation disorder; cancer; cellular adhesion disorder; secreted protein; transmembrane protein; ss.
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                                       Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                  Sequence 18 BP; 2 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                Human PRO298 PCR forward primer 3.
                                                                                     261 ACGGTGCACCTGGAGCA 277
                                                                                                                                                                AAZ34321 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                  98US-0077450P.
98US-0077632P.
98US-0077641P.
98US-0077791P.
98US-0078004P.
98US-0078866P.
                                                                                                          17 Accerecaccareace 1
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                                                                                                                                                                                                            07-DEC-1999 (first entry)
other agents
                                                                                                                                                                                                                                                                                                   Synthetic.
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                    WO9946281-A2
                                                                                                                                                                                                                                                                                                                                                                               08-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                         16-SEP-1999
                                                                                                                                                                                       AAZ34321;
                                                                                                                                           RESULT 835
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                                                                                                                                            Gaps
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145

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This sequence represents a specific example of an antisense oligonucleotide designed to be capable of hybridising to HCV genomic RNA.
These oligonucleotides (AAZ08988-Z09005 and AAZ11701-Z11719) are 10-20 bases long and are targetted to stretches of viral genome which include the polyprotein translation intiation codon. They inhibit the function of viral RNA by interfering with its replication, transcription into protein and packaging into viral particles, resulting in failure of all or a portion of the normal life cycle of the virus. In vivo studies in amvine medel have found that a preferred artisense oligonucleotide, AAZ0893, is able to reduce HCV gene cypression by around 50% compared with a control oligonucleotide are useful for the prevention and/or treatment of hepatitis C-associated disease. The specificity of the oligonucleotides are also useful for detection and diagnosis of hepatitis C virus-associated classase. They can also be used to differentiate between HCV-derived hepatitis and hepatitis caused
                                                                                                                                                                              New antisense oligonucleotides for treatment of Hepatitis C virus
                                                                                                                                                                                                                                                                            Example 3; Page 35; 61pp; English.
                                                        Hanecak RC,
(ISIS-) ISIS PHARM INC.
                                                                                                                     WPI; 1999-493767/41
                                                               Anderson KP,
This sequence represents a specific example of an antisense oligonucleotide designed to be capable of hybridising to HCV genomic RNA.

These oligonucleotides (AAZO1898-Z09005 and AAZO1701-Z1179) are 10-20 bases long and are targetted to stretches of viral genome which include the polyprotein translation initiation codon. They inhibit the function of the polyprotein translation initiation codon. They inhibit the function may, translation into protein and packaging into viral particles, resulting in failure of all or a portion of the normal life cycle of the virus. In vivo studies in a murine model have found that a preferred antisense oligonucleotide, AAZO893, is able to reduce HCV gene expression by around 50% compared with a control oligonucleotide are also useful for detection and diagnosis of hepatitis C virus-associated usease. The specificity of the oligonucleotides more effective prevention and treatment of HCV-derived hepatitis and hepatitis caused the control control and because of the oligonucleotides are also be used to differentiate between HCV-derived hepatitis and hepatitis caused
                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides for treatment of Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 2 A; 4 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                               Nozaki C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 3; Page 35; 61pp; English.
                                                                                          98WO-US026040.
                                                                                                                                                   97US-00988321
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98US-0078936P.
98US-0078939P.
98US-0079656P.
98US-0079663P.
98US-0079663P.
98US-0079786P.
98US-0079786P.
98US-0079788P.
98US-0081237P.
98US-0081237P.
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98US-0081237P.
98US-0081238P.
98US-0081238P.
98US-0081238P.
98US-0081238P.
98US-0081238P.
98US-0081238P.
98US-0081332P.
98US-0081333P.
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22-MAY-1998;
22-MAY-1998;
                                                                                                                                                                                                       31 - MAR - 1998
31 - MAR - 1998
31 - MAR - 1998
01 - APR - 1998
01 - APR - 1998
01 - APR - 1998
08 - APR - 1998
08 - APR - 1998
09 - APR - 1998
09 - APR - 1998
09 - APR - 1998
15 - APR - 1998
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15-APR-1998
21-APR-1998
22-APR-1998
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15-MAY-1998;
15-MAY-1998;
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15-MAY-1998;
15-MAY-1998;
  $\frac{\pi}{\pi} \text{$\frac{\pi}{\pi} \text{$\pi} \text{$\frac{\pi}{\pi} \text{$\frac{\pi}{\pi} \text{$\frac{\pi}{\pi} \text{$\frac{\pi}{\pi} \text{$\frac{\pi}{\pi} \text{$\pi} \text{$\frac{\pi}{\pi} \text{$\pi} \text{$\frac{\pi}{\pi} \text{$\pi} \text{$\frac{\pi}{\pi} \text{$\pi} \text{$\pi}
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The present invention describes secreted and transmembrane polypeptides and their polynucleotides. The nucleotide sequences are useful as sources of probes, primers, for chromosome mapping, and for generation of antisense sequences. They can also be used to create transgenic animals. The proteins can be used to treat a variety of diseases and disorders, depending on their function. Diseases that may be treated include blood cosquilation disorders, cancers and cellular adhesion disorders. They may also be used to raise antibodies. AAZ33891 to AAZ34338, and AAY41685 to AAY41774 represent polynucleotide and polypeptide sequence given in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antisense oligodeoxynucleotide, phosphodiesterase; PDE1B1; enzyme; PDE; cell death; apoptosis; cancer; Ca2+-calmodulin; lymphoblastoid; RNAse H; RPMI 8392; RNA degradation; cAMP; immunoproliferative disorder; breast; immune disfunction; acute lympholytic leukemia; prostate; PCR primer; ss.
                                                                                                                                                                                     New secreted and transmembrane polypeptides and their polymucleotides, useful for treating blood coagulation disorders, cancers and cellular adhesion disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                 Chen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.9%; Score 12.2; DB 1; Length 18; 82.4%; Pred. No. 4.7e+02; vative 0; Mismatches 3; Indels
                                                                                                                                 Baker KP,
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18 BP; 4 A; 6 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                   Yuan J,
                                                                                                                                                                                                                                                      Example 95; Page 257; 530pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human PDE1B1 specific sense primer.
                                                                                                                                   Wood WI, Goddard A, Gurney A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      149 GGAGGCCGGCTTCGACT 165
98US-0086486P.
98US-0087098P.
98US-0087106P.
98US-0087208P.
98US-0094651P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17 GGAGGTCGACTTCCACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX26293 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-MAY-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 2.9
Best Local Similarity 82.4
Matches 14; Conservative
                                                                                                   (GETH ) GENENTECH INC
                                                                                                                                                              WPI; 1999-551358/46.
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22-MAY-1998;
28-MAY-1998;
28-MAY-1998;
28-MAY-1998;
30-UUL-1998;
11-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-SEP-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
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AAX26293
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The invention relates to antisense oligodecxymuclectides (AS-ODM) which will bind to mRNA encoding phosphodiseterase PDEIB1 enzymes and their use in inducting programmed cell death (appotosis) in cancer cells is a Ca2+-calmodulin dependent phosphodiseterase found in cytosolic extracts of human lymphoblastoid cell line, RPMI 8392. The method in which compares the cancer cells (1) applying the phosphodiseterase enzyme PDEIB1 in a cell line containing the cancer cells; (2) synthesising an AS-ODN inhibitor which will bind to the cancer cells; (2) synthesising an AS-ODN inhibitor which will bind to inhibit the enzymatic activity of the PDEIB1 and induce apoptosis in the cells. The AS-ODNs inhibit the expression of a protein by two mechanisms: (1) by degradation of RNA by the ubiquitous enzyme RNase H, which carest of translation initiation caused by AS-ODN bybridization to the S' arrest of translated region or the translation initiation and (ii) the carest of translated region or the translation initiation such expression results in the celevated levels of CAMP. The elevated CAMP levels result in apoptosis by cinhibition of DNA synthesis. The method and AS-ODN are useful in inducing campaign immune disfunctions such as acute lympholytic leukemia,
Antisense oligodeoxynucleotides specific for mRNA encoding phosphodiseterase PBEIB1 enzymes and method for using them to induce apoptosis of cells - useful in the treatment of immunoproliferative disorders and immune disfunctions.
                                                                                                                                                                                                                                                                                    Disclosure; Col 15; 35pp; English
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Natch 2.9%; Score 12.2; DB 1; Length 18; Local Similarity 82.4%; Pred. No. 4.7e+02; Loss 14; Conservative 0; Mismatches 3; Indels Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other; Query Match Best Loca Matches

breast and prostate cancer

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AAX86200 standard; DNA; 18 (first entry) 22-SEP-1999 AAX86200; AAX86200,

ВР

PCR primer used to amplify the PIG3 gene.

p53 transcription tag, p53 status, cancer, cytotoxicity, carcinogenicity, neoplastic, p53 binding site, PIG-3 promoter, PCR primer, ss.

Homo sapiens. Synthetic

WO9914356-A2 25-MAR-1999 98WO-US019300 97US-0059153P. 17-SEP-1998; 17-SEP-1997;

Polyak K; Jogelstein B, Kinzler KW, UNIO) UNIV JOHNS HOPKINS

98US-0079817P.

30-MAR-1998;

WPI; 1999-443793/37.

Use of p53 transcription tags to determine p53 status in, e.g. cancer

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The specification describes the use of p53 transcription tags for developing products to determine p53 status, to diagnose cancer and to evaluate cytotoxicity or carcinogemicity of a test agent. A method for diagnose cancer or determining p53 status in a sample suspected for being neoplastic comprises comparing the level of transcription of an RNA transcription in a first sample (81) of a first tissue (11) to the level of transcription of the transcription a second sample (82) of a second human tissue (62), where 31 is suspected of being neoplastic and 82 is a normal human tissue (of the same type) and the transcription is found to be the same or lower in the first, than in 82. The methods and products can be used to determine p53 status, to diagnose cancer and to evaluate cytotoxicity or carcinogenicity of a test agent.
                                                 Disclosure, Page 14; 73pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                         course of the invention
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Sequence 18 BP; 1 A; 7 C; 7 G; 3 T; 0 U; 0 Other;

Gaps ö Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels Best Loca Matches

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AAX38073 standard; DNA; 18 BP. AAX38073/

RESULT 838

04-JUN-1999 (first entry) AAX38073;

HLA-A specific exon region primer SEQ ID NO:229.

Human; histocompatibility locus antigen; HLA; determination; allele; HLA-B typing; PCR; HLA class I; cis/trans linkage resolution; ss.

Homo sapiens WO9907883-A1 18-FEB-1999

Synthetic

98WO-CA000768. 11-AUG-1998; (VISI-) VISIBLE GENETICS INC. (BLAS/) BLASCZYK R H.

97US-00909290.

11-AUG-1997;

Leushner J; WPI; 1999-167446/14. Blasczyk RH,

Determination of HLA class I group type of a subject - using group specific untranslated region primer pair.

Example, Page 21, 195pp, English.

The present invention describes a method using novel primers involving the PCR-based determination of histocompatibility locus antigen B (HLA-B) class I group type. Determining the HLA-B class I group type of a subject comprises (i) combining a group-specific untranslated region primer pair with a target DNA sample from the subject under conditions such that primer-based amplification of the target DNA may occur; and (ii)

with cell proliferation and inflammation. The antisense oligonucleotides

may also be used as a diagnostic probe for studying gene function

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Sequence 18 BP; 2 A; 5 C; 6 G; 5 T; 0 U; 0 Other;

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The present invention relates to antisense oligonucleotides (see AAA55496 -A55757) which are targeted to nucleic acids encoding a human tumour necrosis factor receptor-associated factor (TRAF). The antisense caquences comprise at least one modified internucleotide linkage, which is a phosphorothicate linkage. The oligonucleotides also include at least one modified supar moiety such as a 2'-0-methoxyethyl sugar moiety. Sequences AAA55490-A55495 represent nucleotide sequences encoding human TRAFI-G. Included in the invention is a method for treating a human having a disease associated with the expression of TRAF comprising administering an antisense oligonucleotide. The reduction of E-cityation in cells comprises comparises comparises comparises contacting the cells or tissues with an antisense oligonucleotide targeted to TRAF-6. In antisense oligonucleotide targeted to TRAF-7 or TRAF-6. The antisense oligonucleotide have antiproliferative and anti-7 or TRAF-6. The antisense oligonucleotide have antiproliferative and anti-7 or TRAF-6.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense oligonucleotides targeted to nucleic acids encoding human tumor necrosis factor receptor-associated factor (TRAF), useful for treating diseases associated with TRAF expression such as inflammatory diseases.
              amplification; where the ability of the primer pair to produce a nucleic acid product is associated with a particular HIA group type. The method can be used for HIA-B typing. In the method, the initial group specific amplification allows a PCR based separation of haplotypes in 95% of patient samples. It permits the resolution of cis/trans linkages of heterozypote sequencing results which cannot be achieved with other protocols. AAX37845 to AAX3886 represent DNA sequence used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tumour necrosis factor receptor-associated factor; TRAF; human; antisense oligonucleotide; phosphorothioate; antiproliferative; anti-inflammatory; E-selectin; jun kinase; ss.
                                                                                                                                                                                                                                                                                                             ö
determining whether a nucleic acid product is produced by the
                                                                                                                                                                                                                                                               2.9%; Score 12.2; DB 1; Length 18; 82.4%; Pred. No. 4.7e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                       Seguence 18 BP; 1 A; 7 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TRAF1 antisense oligonucleotide ISIS# 26707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Monia BP, Xu XS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 14; Page 46; 170pp; English
                                                                                                                                                                                                                                                                                                                                                        214 AGAACTCGGTGGCGCC 230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAASSSOS standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99WO-US023171.
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                                                                                                                                                                                                                                                                                                                                                                                  18 ACAACTGGGAGGCGGCC 2
                                                                                                                                                                                                                                                                                       ilarity 82.4%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-303732/26.
                                                                                                                                                                                                                                                                                       Local Similarity
Les 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200020435-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA55505;
                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 839
                                                                                                                                                                                                                                                                                            Best Loca
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                   Gaps
                                                                                                                                                                                                                                                                                                                  Tumour necrosis factor receptor type 1; TNFR1; antisense; infection; inflammation; tumour formation; TNFR1; anticancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense inhibition of tumor necrosis factor type 1 expression for diagnosis, treatment and prevention of disease, particularly tumors
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2.9%; Score 12.2; DB 1; Length 18;
llarity 82.4%; Pred. No. 4.7e+02;
Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                     Human TNFR1 mRNA inhibiting antisense oligo ISIS# 18941.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18 BP; 3 A; 9 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Col 25; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               131 GCTGGCCCGCCTGGCGG 147
                                                                   239 AGGCTGCTTCCCGGGCT 255
                                                                                                                                                                                   AAZ48548 standard; DNA; 18 BP
                                                                                                1 AGACGGCTTCCTGGGCT 17
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                                                                                                                                                                                                                                                       31-MAR-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ISIS-) ISIS PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-105333/09.
                 Local Similarity
hes 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
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     Query Match
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AAZ39609
                     Best Loca
Matches
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Seguence 18 BP; 2 A; 3 C; 6 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                                                                             Chumakov I;
                                                                                                                                                                                                                                                                                                                                                              Claim 8; Page 1125; 2745pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAC78898 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18 GACCACCACTTAGAGAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18-FEB-2000; 2000WO-US004341
                                                                                          99WO-IB000822.
                                                                                                                             98US-0082614P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-FEB-2001 (first entry)
                                                                                                                                                                                                                         Cohen D, Blumenfeld M,
                                                                                                                                                                                                                                                                WPI; 2000-013267/01.
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                                                                                                                                                                                      (GEST ) GENSET.
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                                                                                                                               21-APR-1998;
23-NOV-1998;
                WO9954500-A2
                                                                                          21-APR-1999;
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                                                     28-OCT-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention provides antisense compounds targeted to a coding region, yirk or 5 urgs of a nucleic acid molecule encoding human cREL (transcriptional activator). The antisense compounds are useful as research agents and diagnostics such as in the elucidation of the thortion of a particular gene. The antisense compounds can be useful as therapeutic modalities that can be configured to be useful in treatment regimes for treatment of cells, tissues and animals, especially humans. In the prior art, there are no known therapeutic agents which effectively inhibit the synthesis of cREL and additional agents capable of inhibiting CREL function are still required. Sequences AAZ39589-627 represent
                                                                                                                                                 Human; cREL; transcriptional activator; antisense compound; therapeutic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antisense phosphorothioate oligodeoxynucleotides inhibiting human CREL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
2.9%; Score 12.2; DB 1; Length 18;
Best Local Similarity 82.4%; Pred. No. 4.7e+02;
Matches 14; Conservative 0; Mismatches 3; Indels 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human biallelic marker upstream amplification primer SEQ ID NO:4194.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human genome, biallelic marker, high density disequilibrium map, genomic map, haplotype, phenotype, polymorphic base, genotyping, haplotyping, hybridisation, identification, characterisation, amplification, single nucleotide polymorphism; SNP; PCR primer,
                                                                                                            Human CREL mRNA inhibiting antisense oligo ISIS #24093.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18 BP; 2 A; 5 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense modulation of human cREL expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 15; Col 27; 26pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Baker BF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TTCTACGTGATCGAGAC 421
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AAZ39609 standard; DNA; 18 BP.
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                                                                                                                                                                                                                                                                                                                                              98US-00156253
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                                                                         28-FEB-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Monia BP, Cowsert LM,
                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-061889/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               diagnosis; ss.
                                                                                                                                                                                                                             Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                              18-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                     18-SEP-1998;
                                                                                                                                                                                                                                                                  US6001652-A
                                                                                                                                                                                                                                                                                                       14-DEC-1999
                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          405
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AAZ69838/C
XX
AAC AAZ6983
XX
DT 10-SEP.
XX
XX
HUMAN |
XX
KW HUMAN |
XW ABOOM!
KW AMPIOT
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Homo sapiens

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CC invention, which contain a polymorphic base at position 24 of their invention, which contain a polymorphic base at position 24 of their contedide sequences. AAZ65579 to AAZ7740 reppresent amplification contain a polymorphic base at position 24 of their contests. AAZ65791 to AAZ7740 reppresent amplification primers for the biallelic markers. The biallelic markers of the invention primers for the biallelic markers. The biallelic markers of the invention contains and markers association studies and haplotyping studies which are useful in determining the genetic basis for disease states. CC compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents acting on a disease to and side effects from the sequence contains acting on a disease as well as other treatment.

N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3056, 3157, 3227, 3297 and 3567, are not actually given a sequence in the Sequence Listing from the contains and the contains
Novel biallelic markers used to construct a high density disequilibrium map of the human genome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human, secreted protein, transmembrane protein, PRO, BST, cytostatic, expressed sequence tag, detection, cancer, PCR primer, probe, 8s.
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99US-0123957P.
99US-0126773P.
99US-0130232P.
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29-MAR-1999;
21-APR-1999;
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DE NEMOURS & CO

99US-0120702P

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a large number of arbitrarily primed polymerase chain reactions comprises separating two populations of microbial cells, where a first population is contacted with a stimulating agent, extracting total RNA from both microbial cells pupulation applicyly application of at least thirty-two both contacted with a stimulating agent, extracting total RNA from both microbial cell populations, amplifying the extracted RNA from both populations by preparing a collection of at least thirty-two different arbitrary primers, where each primer comprises a common and a variable region; individually contacting each primer of with a sample of extracted region; individually contacting each primer of with a sample of extracted amplification products are produced; purifying the two sets of amplification products are produced; purifying the two sets of amplification products are produced; purifying the two sets of amplification which differ from products generated from the second population which differ from products generated from the second population which differ from products generated from the equencing the identified differentially expressed genes. The advantage of one genes using thirty-two or thirty primers have isolated four and one clone genes using thirty-two or thirty primers have isolated four and one clone genes using thirty-two or thirty primers have isolated four and one clone genes using thirty-two or thirty primers have isolated twenty-one induced gene fragments. This universal primers has isolated twenty-one induced gene fragments. This universal primer was used for the reamplification of the differentially amplified bands
                                                                                                                                                                                                                                                                                                                                                                                                                     Differential display method using a large number of arbitrary primers for RT-PCR used to isolate novel differentially expressed prokaryotic genes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure, Page 61; 66pp; English
                                                                         17-FEB-2000; 2000WO-US003989
                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-587069/55.
                                                                                                                                                                                                              TNOW DU ( OGUU)
                                                                                                                                                   19-FEB-1999;
                                                                                                                                                                                                                                                                                        Rouviere P;
           8X#X#X#X#X#X##X##X##X#X#X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
Goddard A, Godowski PJ, Grimaldi CJ, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL;
Stewart TA, Tumas D, Williams PM, Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel PRO polypeptides and polynucleotides used in detection methods, t
target bioactive molecules to specific cells, and to modulate cellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Seguence 18 BP; 4 A; 6 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 95; Page 317; 636pp; English.
28-APR-1999; 99US-0131445P.
14-MAY-1999; 99US-0134287P.
26-UUL-1999; 99US-0141037P.
26-UUL-1999; 99US-0145698P.
20-OCT-1999; 99WO-USC02813.
02-DEC-1999; 99WO-USC028551.
02-DEC-1999; 99WO-USC028565.
16-DEC-1999; 99WO-USC031243.
30-DEC-1999; 99WO-USC031243.
30-DEC-1999; 99WO-USC031243.
30-DEC-1999; 99WO-USC031274.
06-JAN-2000; 2000WO-USC00277.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                VPI; 2000-611443/58.
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Motilin receptor; gastrointestinal disease; gastric motility disorder;
gastroparesis; irritable bowel syndrome; diarrhoea; ss.
                                                                                                                                                                                                        Polynucleotide in unique region in exon 1 of rabbit motilin receptor
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Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                  178 AGTCCAAGGCACATATC 194
                                                                 1 AGTCCACGGAGCATATC 17
                                                                                                                                    AAF85457 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                   29-OCT-1999; 99US-0162264P.
                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                          (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                      Oryctolagus cuniculus.
                                                                                                                                                                                                                                                                                             WO200132710-A1
                                                                                                                                                                                   23-JUL-2001
                                                                                                                                                                                                                                                                                                                     10-MAY-2001
                                                                                                                                                             AAF85457;
                                                                                                             RESULT 845
AAF85457/c
                                                                                                                                     ઠે
                                                                         셤
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Identification; prokaryote; polymerase chain reaction; PCR; amplification; primer; differential display; picric acid degredation; agene cluster; open reading frame; ORF; dehydratase; dehydrogenase; transcription factor; Acyl-CoA synthase; NADPH oxidoreductase; ss.

WO200049177-A2

Synthetic

Universal primer used in differentiation/identification method

(first entry)

03-JAN-2001

AAA53953;

BP.

AAA53953 standard; DNA; 18

RESULT 844 AAA53953

Ö

0; Gaps

Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels

149 GGAGGCCGGCTTCGACT 165

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17 deaderceacriceacr 1

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29-DEC-1999; US6187586-B1 Query Match 846 AAF79645/ RESULT ઠે 셤

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AAF85456-60 represent polynucleotide sequences from the unique region of exon 1 of a rabbit motilin receptor gene. The specification describes an unique sequence present in exon 1 of the motilin receptor, which is not present in human or Sphaeroides nephalus 75E7 motilin receptor sequences. The unique nucleic acid sequence is useful for measuring the ability of a compound to affect motilin receptor activity. Motilin receptor compound to affect motilin receptor activity. Motilin receptor compounds which are useful for treating gastrointestinal diseases and disorders such as gastric motility disorders, gastroparesis, irritable bowel syndrome, and diarrhoea
                                                                                                                                                                                          Novel polypeptides related to dog and rabbit motilin receptor polypeptide, comprising unique regions from dog and motilin receptor amino acid sequence, useful for identifying compounds for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 1 A; 7 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 17; Page 22; 42pp; English.
                                                                                          WPI; 2001-343479/36
                                                                                                                                                                                                                                                                                                                                          diarrhea in humans.
Mckee K;
Tan C,
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2.9%; Score 12.2; DB 1; Length 18; 82.4%; Pred. No. 4.7e+02; Aative 0; Mismatches 3; Indels
                                                                                 373 TCCTGGACCGCGACGAC 389
                                                                                                                       N
                                                                                                                       TCCGGGGCCGCGAAGAC
                       Local Similarity 82.4
les 14; Conservative
                                                                                                                         18
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Human Akt-3 antisense oligonucleotide, SEQ ID NO: 53.
ВР
AAF79645 standard; DNA; 18
                                                                         (first entry)
                                                                         29-MAY-2001
                                      AAF79645;
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Human, Akt-3; protein kinase; cytostatic; antiinflammatory; infection; antisense therapy; inflammation; tumour; ss.

Homo sapiens

13-FEB-2001.

99US-00474922 29-DEC-1999; 99US-00474922. (ISIS-) ISIS PHARM INC. Roth RA; Cowsert LM, Monia BP,

WPI; 2001-264979/27.

The present sequence is one of a number of antisense compounds of up to 30 nucleobases in length targeted to a nucleic acid encoding human Akt-3. The antisense compounds are useful for inhibiting the expression of human Akt-3 in human cells or tissues. They are also useful for modulating the expression of Akt-3, and for treating a human or an animal suspected of New antisense compounds targeting nucleic acids encoding human Akt-3 useful for treating a disease or condition associated with Akt-3 expression, or in preventing or delaying inflammation or tumor formation. Claim 1; Col 39; 37pp; English.

having, or being prone to, a disease or condition associated with Akt-3 expression. The antisense compounds may also be used as research reagents, in kits and in diagnostics, e.g. to elucidate the function of particular gene or to distinguish between functions of various members ca biological pathway; and as a prophylactic, e.g. to prevent or delay infection, inflammation or tumour formation

8888888

Sequence 18 BP; 3 A; 6 C; 4 G; 5 T; 0 U; 0 Other;

ö Gaps ö 2.9%; Score 12.2; DB 1; Length 18; 82.4%; Pred. No. 4.7e+02; tive 0; Mismatches 3; Indels Local Similarity 82.4 nes 14; Conservative Query Match Matches

92 N 76 AGGGCCGCGCAGTGGAC 18 Ardecedadeadradae

g à

ВР

RESULT 847

Rat P00188D09 RNA reverse PCR primer.

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Gaps

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Rat; secreted factor; P00210D09; P0018BD09; cardiant; nephrotropic; antiinflammatory; gene therapy; cardiac disease; renal disease; inflammatory disease; PCR primer; ss.

27-SEP-2000; 2000WO-US026582.

99US-0156277P

Novel secreted factor encoded by clone F00210D09 useful for diagnosing, treating and/or preventing various cardiac, renal and inflammatory

Example 9; Page 51; 69pp; English.

The present sequence was used to amplify rat P00188D09 RNA by quantitative real-time PCR. The invention relates to a polypeptide comprising a sequence of at least 80% identity to residues 22-122 of the present sequence, or a sequence encoded by a nucleic acid hybridising under stringent conditions to the complement of the coding region comprising 1031 nucleotides, and having at least one biological activity of the polypeptide encoded by rat clone p00210D09. The polypeptides and polymucleotides of the invention are useful for the treatment of cardiac, renal and inflammatory diseases. The P0021D09 polymucleotides are useful in antisense mediated gene inhibition and in gene therapy. The polypeptides are useful in assays for identifying lead compounds that may be used as therapoutic agents in the treatment of cardiac, is and in the treatment of cardiac, is a sequence of the compounds that may be asset in the treatment of cardiac, kidney or

Seguence 18 BP; 3 A; 7 C; 4 G; 4 T; 0 U; 0 Other;

Gaps .. 0 2.9%; Score 12.2; DB 1; Length 18; larity 82.4%; Pred. No. 4.7e+02; Conservative 0; Mismatches 3; Indels Best Local Similarity Matches 14; Conserv Query Match

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16G ADT 21 12:58:21 2004

TGCGGGTGACCGAGGGC 32 ||| ||||| |||| 17 TGCAGGTGATCGACGGC 1

ò d

AAH40381 standard; DNA; 18 BP.

AAH40381;

14-AUG-2001 (first entry)

SNP specific upper PCR primer SEQ ID 3177.

Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipldus; cancer; lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyxia; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.

Homo sapiens

40200129262-A2.

26-APR-2001.

13-OCT-2000; 2000WO-US028436.

.5-OCT-1999; 99US-0160096P.

ORCH-) ORCHID BIOSCIENCES INC.

Picoult-Newburg L, Pohl M;

VPI; 2001-290930/30.

New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic acid sample.

Claim 1; Page 66; 83pp; English.

Sequences AAH17205 - AAH40944 represent PCR primers, single nucleotide polymorphisms SNPs. The present invention is ties of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a CSNP inalking sequence, the SNPs primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The coligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. or agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, familial hypercholesteroleamia, polycystic. Phenotypic traits also include symptoms of or susceptibility to multifactorial diseases of which a component is or may be genetic such as autoimmune diseases, including, rheumatoid arthritis, multiple sclerosis, including, rheumatoid arthritis, multiple sclerosis, incroorganism. The method is also useful in forensic investigations and paranty analysis. The present represents a PCR primer specific for a human SNP containing DNA sequence

Sequence 18 BP; 4 A; 4 C; 9 G; 1 T; 0 U; 0 Other;

2.9%; Score 12.2; DB 1; Length 18; 82.4%; Pred. No. 4.7e+02; Query Match Best Local Similarity

Indels ы ., Mismatches ö Matches 14; Conservative

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156 140 CCTGGCGGTGGAGGCCG 2 ccederaderenandecce

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ABZ72355/c ID ABZ72355 standard; DNA; 18

ABZ72355;

(first entry) 03-APR-2003 Gene 216 polymorphism genotyping ASO primer SEQ ID NO 327.

Human, Gene 216; chromosome 20p13-p12; antiasthmatic; anorectic; antiinflammatory; gastrointestinal; gene therapy; vaccine; asthma; obesity; inflammatory bowel disease; primer; ss.

Synthetic.

WO200178894-A2.

25-OCT-2001.

13-APR-2001; 2001WO-US012245.

(GENO-) GENOME THERAPEUTICS CORP.

13-APR-2000; 2000US-00548797.

Keith T;

WPI; 2001-639428/73.

Isolated genes (Gene 216) from human chromosome 20p13-p12 and the proteins they encode, useful for the prevention, diagnosis and treatment of asthma, obesity and inflammatory bowel disease.

Example 11; Page 156; 520pp; English.

The invention relates to isolated genes (Gene 216) from human chromosome 20p13-p12 and the proteins they encode. The mucleic acids and proteins may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate Gene 216 expression. For example, the nucleic acids (or vectors) and proteins may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of gene 216 by expressing inactive proteins or to supplement the patients own production of Gene contractions. Additionally, the nucleic acids may be used to produce the sereted Gene 216 protein, by inserting the nucleic acids into a host complementary sequences may also be used as DNA probes in diagnostic coll and culturing the cell to express the protein. The nucleic acids and collect acids and collect acids and contraction of antibodies and therefore which patients may be in need of restorative therapy. The Gene 216 protein may also be used as antigens in capturity. The anti-Gene 216 expression and activity. The proteins may also be used as diagnostic agents for dececting the presence of Gene 216 proteins in samples (e.g. by enzyme linked immunosorbant assay or ELISA). Disorders that may be conformation. The primers are used in the physical mapping of the gene C Abazzoes), polymorphism (dentification using single strand conformational polymorphism (SSCP) analysis (ABZ720114-ABZ72184), credenting (ABZ72189), and genetyping (ABZ721191-ABZ72184).

Sequence 18 BP; 1 A; 3 C; 9 G; 5 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mass (HBM) gene, which are found on chromosome 11q13.3. The ZmaxI and HBM genes have osteopathic activities. The genes can be used in gene therapy, antisense therapy and in the production of vaccines. They can be used in the diagnosis and treatment of bone disorders including osteoporosis, Pager's disease, sclerostosis, osteomalacia and fibrous dysplasia. ABA82038 to ABA82700 and AAG68168 to AAG68193 represent sequences used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes the human Zmax1 gene and the high bone
                                                                                                                                                                                                                                                                                                                      Human; high bone mass; HBM gene; Zmaxl gene; chromosome 11; 11q13.3; sequence tagged site; STS; osteopotals; osteopathic; gene therapy; antisense therapy; vaccine; bone disorder; Paget's disease; adapter; sclerostosis; osteomalacia; fibrous dysplasia; PCR primer; linker; ss.
                                  Gaps
                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New high bone mass (HBM) and Zmaxl genes and proteins useful modulating bone mass for the treatment of e.g. osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 12.2; DB 1; Length 18; Pred. No. 4.7e+02; 0; Mismatches 3; Indels
Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                         gene region physical map preparation STS marker #235.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seguence 18 BP; 3 A; 8 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Carulli JP, Little RD, Recker RR, Johnson ML,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 34; 443pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GENO-) GENOME THERAPEUTICS CORP.
                                                                  CATCACCACGTCTGACC 108
                                                                                                                                                                                       ВР.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-APR-2000; 2000US-0054371.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-JUN-2000; 2000WO-US016951
                                                                                               CAGCACCACAGCTGACC
                                                                                                                                                                                    ABA82276 standard; DNA; 18
                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-657171/75
                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200177327-A1.
                                                                                                                                                                                                                                                     25-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                  92
                                                                                               17
                                                                                                                                                                                                                    ABA82276;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                         Zmax1
                                                                                                                                                     RESULT 850
                                                                                                                                                                                                                                                                                                                                                                                                              Ношон
                                                                                                                                                                         ABA82276
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ö ö Length 18; 3; Indels Sequence 18 BP; 1 A; 8 C; 5 G; 4 T; 0 U; 0 Other; 2.9%; Score 12.2; DB 1; 82.4%; Pred. No. 4.7e+02; iive 0; Mismatches 3; 14; Conservative Query Match Best Local Similarity

1 N GAGGGCCGCGCAGTGGA GATGGCCCCGCAGAGGA 75 18 용 δ

Gaps

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Local Similarity 82.4

Matches

ઠે g

1 GGCCAGGAGTGAAACTG 17

descadeadraacrerd

ABT05044 standard; DNA; 18 ABT05044; RESULT 852
ABT05044/C
ID ABT0504
XX
AC ABT0504

ВP

AAS20963 standard; DNA; 18

AAS20963,

AAS20963

09-APR-2002 (first entry)

PCR primer Igf2r-12 relating to gene imprinting invention.

Human; genomic imprinting; pluripotent mouse embryonic germ cell line; BG; methylated CpG island; DNA methylation; gene imprinting; post-translational modification of histone; cancer; birth defect; diabetes; aberrant imprinting; PCR; primer; ss.

Homo sapiens

WO200190313-A2.

29-NOV-2001.

22-MAY-2001; 2001WO-US016253

2000US-0206158P. 2000US-0206161P. 22-MAY-2000; 22-MAY-2000;

(UYJO) UNIV JOHNS HOPKINS.

Strichman-Almashanu L, Jiang Feinberg A,

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WPI; 2002-083100/11.

Forming embryonic germ cells useful as model system to study imprinting involves mating genetically divergent male and female mammal of same species, dissecting and dissociating embryo obtained from pregnant

Disclosure; Page 54; 125pp; English.

The present invention relates to a model system for genomic imprinting using pluripotent mouse embryonic germ (BG) cell lines derived from an interpretific cross. Also disclosed is a library containing methylated confidence and a method for assaying methylation in one or more imprintable genes. The gene imprinting assay is carried out by single or strength on polymorphism (SSCP), quantitative sequencing, single nucleotide primer extension or hot stop PGR. The assays are carried out to determine the post-translational modification of histones. The method to determine the post-translational modification of histones. The method to determine the post-translational modification of histones. The method to determine the post-translational modification of histones. The method of contracting cancer if the test substance enhances imprinting of a gene whose imprinting is gained in cancer. The methylated CpG islands are useful for providing diagnostic information relative to developing cancer, or for providing diagnostic information relative to cancer which involves determining the methylation status of the risk of elemes which are useful for detecting birth defects, diabetes and cancers assaciated with aberrant imprinting. The EG cell lines represent the first in vitro model system in which genomic imprinting can be followed dynamically and the two alleles can be distringuished. AAS20953-AAS20969 represent PCR primers described in the present invention

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nse compound; tumour necrosis factor receptor 1; liver disease; hepatitis; liver injury; hyperproliferative disorder; cancer; ds.
          INFR1 expression modulation related antisense oligo SEQ ID No 74.
                                                                         22-OCT-2001; 2001WO-US051224.
                                                    WO200248168-A1.
                                           Homo sapiens.
11-OCT-2002
                                                              20-JUN-2002
                     Antisense
                                human;
```

24-OCT-2000; 2000US-00695451.

'n Zhang Cowsert LM, (ISIS-) ISIS PHARM INC Baker BF,

WPI; 2002-583481/62

Dean NM;

Novel antisense compound targeted to nucleic acid molecule encoding tumor necrosis factor receptor 1 (TNPR1), useful for treating humans having disease associated with TNFR1 e.g. hepatitis, liver injury, liver cancer.

Example 10; Page 45; 121pp; English

The invention relates to an antisense compound 8 to 30 nucleotides in length targeted to nucleic acid molecule encoding tumour necrosis factor receptor 1 (TNFR1), where the antisense compound inhibits expression of TNFR1. The antisense compound is useful for inhibiting the expression of Treating an animal (preferably human) having a disease or condition associated with TNFR1, e.g. a liver disease (such as hepatitis, or liver injury) or a hyperproliferative disorder such as cancer, by inhibiting the expression of TNFR1. The antisense compound is useful for injury or a hyperproliferative disorder such as cancer, by inhibiting diagnostics, therapeutics, prophylaxis and as research reagents and kits. This polynucleotide sequence represents a human oligonucleotide relating to the TNFR1 of the invention

Sequence 18 BP; 3 A; 9 C; 4 G; 2 T; 0 U; 0 Other;

0; Gaps ch 2.9%; Score 12.2; DB 1; Length 18; 1 Similarity 82.4%; Pred. No. 4.7e+02; 14; Conservative 0; Mismatches 3; Indels Query Match Best Local 9

131 GCTGGCCCGCCTGGCGG 147 GCTGGGCTGCCTGGAGG 18

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ABT05119 standard; DNA; 18 BP.

11-OCT-2002 ABT05119;

(first entry)

INFR1 expression modulation related antisense oligo SEQ ID No 149

Antisense compound; tumour necrosis factor receptor 1; liver disease; TNFR1; hepatitis; liver injury; hyperproliferative disorder; cancer; human; ds.

Homo sapiens

40200248168-A1

20-JUN-2002

22-OCT-2001; 2001WO-US051224

24-OCT-2000; 2000US-00695451

(ISIS-) ISIS PHARM INC

Dean NM Zhang H, Cowsert LM, Baker BF,

WPI; 2002-583481/62

Novel antisense compound targeted to nucleic acid molecule encoding tumor necrosis factor receptor 1 (TNFR1), useful for treating humans having disease associated with TNFR1 e.g. hepatitis, liver injury, liver cancer.

Example 18; Page 56; 121pp; English

The invention relates to an antisense compound 8 to 30 nucleotides in length targeted to nucleic acid molecule encoding tumour necrosis factor receptor 1 (TNRM1), where the antisense compound inhibits expression of TNRM1. The antisense compound is useful for inhibiting the expression of TNRM1 in cells or tissues. The antisense compound is also useful for treating an animal (preferably human) having a disease or condition associated with TNRM1, e.g. a liver disease (such as hepatitis, or liver injury) or a hyperproliferative disorder such as cancer, by inhibiting the expression of TNRM1. The antisense compound is useful for diagnostics, therapeutics, prophylaxis and as research reagents and kits. This polynucleotide sequence represents a human oligonucleotide relating to the TNRM1 of the invention

Sequence 18 BP; 4 A; 2 C; 9 G; 3 T; 0 U; 0 Other;

Gaps ö 2.9%; Score 12.2; DB 1; Length 18; 32.4%; Pred. No. 4.7e+02; ve 0; Mismatches 3; Indels Query Match Best Local Similarity 82.4%; Matches 14; Conservative

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ò g RESULT 854 AAL43633 ID AAL4

AAL43633 standard; DNA; 18 BP.

AAL43633;

Rhodococcus picric acid degradation pathway-related universal PCR primer. (first entry) 05-SEP-2002

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Picric acid degradation gene cluster; ss; recombinant organism; picric acid degradation pathway; PCR; primer.

Unidentified.

US2002042117-A1.

11-APR-2002.

03-SEP-1999; 99US-0152545P. 31-AUG-2000; 2000US-00651941. 17-SEP-2001; 2001US-00955597

(ROUV/) ROUVIERE P.E. (WALT/) WALTERS D.M. (RUSS/) RUSS R.

ä Russ Rouviere PE, Walters DM,

WPI; 2002-381946/41

Isolated nucleic acid fragments encoding enzymes of the picric acid degradation pathway isolated from Rhodococcus erythropolis HL PM-1, useful in the creation of recombinant organisms that have the ability picric acid. degrade

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Example 5; Page 15; 53pp; English

The invention comprises 12 Rhodococcus erythropolis ORFs encoding enzymes of the picric acid degradation pathway. The invention also comprises the nuclectide sequence of the picric acid degradation gene cluster containing all 12 of the ORFs. The picric acid degradation pathway genes and enzymes of the invention are useful for creating recombinant organisms that have the ability to degrade picric acid. As well as for the identification of new species of bacteria that have the ability to degrade picric acid. The picric of new species of bacteria that have the ability to degrade picric acid. The present DNA sequence represents a Rhodococcus picric acid degradation pathway-related universal reamplification PCR primer

Sequence 18 BP; 5 A; 5 C; 5 G; 3 T; 0 U; 0 Other;

Gaps .. 0 2.9%; Score 12.2; DB 1; Length 18; 82.4%; Pred. No. 4.7e+02; ative 0; Mismatches 3; Indels 194 Best Local Similarity 82.4 Matches 14; Conservative Query Match ò g

ABK51851 standard; DNA; 18 (first entry) 13-AUG-2002 ABK51851; ABK5185 RESULT

BP

Picric acid degradation; 2,4,6-trinitrophenol; explosive manufacturing; aniline; colour fast dye; pharmaceutical; steel etching; PCR; environmental toxicant; enzymatic degradative process; primer; ss. R. erythropolis picric acid degradation related universal PCR primer.

Synthetic.

US6355470-B1.

12-MAR-2002.

31-AUG-2000; 2000US-00651941

99US-0152545P 03-SEP-1999; DUPO) DU PONT DE NEMOURS & CO E I.

Rouviere PE, Walters DM, Russ

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WPI; 2002-433274/46.

Nucleic acid encoding an F420/NADPH oxidoreductase isolated from Rhodococcus erythropolis HL PM-1 is associated with picric acid degradation and is useful to create recombinant organisms that degrade. Example 5; Col 26; 49pp; English

The present invention relates to the isolation of Rhodococcus erythropolis Hi PM-1 gene cluster containing 12 open reading frames (ORFs) implicated in the degradation of picric acid (2,4,6-1) trinitrophenol). The polynuclectide sequences of the invention are useful for creating recombinant organisms that have the ability to degrade picric acid. Ploric acid which is used in industrial applications including the manufacture of explosives, aniline, colour fast dyes, pharmaceuticals and in steel etching, is highly unstable. The present

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invention provides a means of disposal/removal of this toxic substance from the environment by an enzymatic degradative process. The present sequence represents a universal PCR primer used in the examples of the
                                                                      present invention
       8888888
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Sequence 18 BP; 5 A; 5 C; 5 G; 3 T; 0 U; 0 Other;

Gaps . 0 Length 18; 3; Indels Score 12.2; DB 1; Pred. No. 4.7e+02; 0; Mismatches 2.9%; Query Match 2.9 Best Local Similarity 82.4 Matches 14; Conservative

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ABK23073 standard; DNA; 18 RESULT 85 ABK23073/ ID ABK2

ВР

ABK23073;

(first entry) 09-APR-2002

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Human Zmax1 cDNA forward PCR primer #118.

Human; mouse; Zmax1; HBM; high bone mass gene; lipid regulation; stroke; lipid-associated condition; arteriosclerosis; cardiovascular disease; 85; osteoporosis; atherosclerosis; diabetic atherosclerosis; plaque build-up; neurovascular condition; wound healing; gene therapy; PCR primer; probe; bone development disorder; antiarteriosclerotic; cardiovascular; osteopathic; cerebroprotective.

Homo sapiens

WO200192891-A2.

06-DEC-2001.

25-MAY-2001; 2001WO-US016946.

26-MAY-2000; 2000US-00578900.

(GENO-) GENOME THERAPEUTICS CORP. (UYCR-) UNIV CREIGHTON SCHOOL MEDICINE.

Carulli JP, Little RD, Recker RR, Johnson ML;

WPI; 2002-097784/13.

Identifying molecules involved in lipid regulation, useful for diagnosing, treating or preventing e.g., arteriosclerosis, comprises identifying a molecule that binds to high bone mass gene or its corresponding wild type gene.

Disclosure; Page 39; 409pp; English.

The invention relates to a method for identifying a molecule involved in lipid regulation comprising identifying a molecule that binds to or comprising identifying a molecule that binds to or inhibits binding of a molecule to high bone mass (HBM) or its wild type gene, Zmax1. Compounds identified by the method are useful for treating, diagnosing, preventing or screening for normal and abnormal lipid associated conditions, including arteriosclerosis, cardiovascular fractment or prevention of diabetic atherosclerosis, neurovascular conditions caused by plaque build-up, poor circulation due to plaque build-up, poor circulation due to plaque conditions dassociated poor wound healing. The methods may be used in conditions as associated poor wound healing. The methods may be used in care therapy, pharmaceutical development, and diagnostic assays for bone development, in diagnosis of human or animal bone disease, and in the treatment of bone diseases. Sequences ABK22776-ABK23411 represent cDNA molecules encoding human Zmax1 and HBM, and PCR primers, probes, linkers

Human Her-2 antisense oligonucleotide, ISIS #27972.

(first entry)

23-SEP-2002

AAD38945;

AAD38945 standard; DNA; 18 BP.

4AD38945

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The invention relates to a typing kit for judging human leukocyte antigen (HLA) genocype of a sample by hybridising a substrate on which 10-24 base oligonucleotides (AB130512-AB131809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals
                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when transplanting between them.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nishida M;
                                                                                                  Gaps
                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                               Human, human leukocyte antigen, HLA, genotype, polymorphism,
immunogenetic, transplantation, genetic disease, ss.
                                                                   Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matsumura Y, Moriya S,
                                                                                                3, Indels
                                                                                                                                                                                                                                                                                                                                                                 Human HLA genotyping oligonucleotide SEQ ID NO 187.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 4 A; 4 C; 7 G; 3 T; 0 U; 0 Other;
                               Sequence 18 BP; 3 A; 8 C; 3 G; 4 T; 0 U; 0 Other;
                                                             Query Match
2.9%; Score 12.2; DB 1;
Best Local Similarity 82.4%; Pred. No. 4.7e+02;
Matches 14; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 10; Page 128; 345pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ichihara T,
                                                                                                                                                                                                                                                             ABL30698 standard; DNA; 18 BP.
and adapters of the invention
                                                                                                                                      1 GGCCAGGAGTGAACTG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-JUN-2000; 2000JP-00164798.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-JUN-2001; 2001WO-JP004662.
                                                                                                                                                                 18 GGCAGAGTGACTCTG 2
                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (NISN ) NISSHINBO IND INC. (SYST-) SYSTEM RES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Inoko H, Kagiya T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-122074/16.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                21-MAR-2002
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                                                                                                                                                                                                                                                                                            ABL30698;
                                                                                                                                                                                                                             857
                                                                                                                                                                                                                           RESULT 85
ABL30698
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The invention relates to antisense compounds targetted to a nucleic acid molecule encoding Her2 (human Epidermal Growth Factor receptor 2) that specifically hybridises with and inhibits the expression of Her2. Antisense compounds of the invention are used for treating diseases or conditions associated with Her2 such as hyperproliferative disorders e.g. lung, breast, gastric, oesophageal, colon, bladder, salivary, neural or cardiac cancer. They are also useful prophylactically e.g. to prevent or delay infection, inflammation and tumour formation. The invention is also used in gene therapy. The present sequence is an antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel antisense oligonucleotide which modulates the expression of Huma
Epidermal Growth Factor receptor, Her2, is useful for treating tumors
inflammation or to prevent infection in humans
                                                                                          Human; Her-2; epidermal growth factor receptor 2; infection; cancer; hyperproliferative disorder; prophylaxis; inflammation; antisense; tumour; gene therapy; phosphorothicate backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= m5c
15. .18
/*tabas c
/mod_base OTHER
/note= "2'methoxyethyl nucleotides"
                                                                                                                                                                                                                                                     1. .4
/*tea b
//mod base= OTHER
/note= "2'methoxyethyl nucleotides"
                                                                                                                                                                                                 1..18
/*tag= a
/mod bass= OTHER
/note= "Phosphorothioate backbone"
                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 89; 116pp; English.
                                                                                                                                                                                                                                                                                                                                                                 mod_base= m5c
                                                                                                                                                                                                                                                                                                                              /mod_base= m5c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-SEP-2000; 2000US-00663834.
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                                                                                                                                                                                                                                                                                                                                                                                            tag=
                                                                                                                                                                                                                                                                                                                  *tag=
                                                                                                                                                                                                                                                                                                                                                       *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ISIS-) ISIS PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200222636-A1
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modified_base
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                                                                                                                                                Homo sapiens
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-MAR-2002
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Sequence 18 BP; 4 A; 3 C; 9 G; 2 T; 0 U; 0 Other;

Gaps

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Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels

6 g primer.

BP.

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Neublastin DNA related PCR
                                                                         ABT11916 standard; DNA; 18
                                                                                                                                                                                                              19-DEC-2002
                                                                                                                                                ABT11916;
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RESULT B ABT 11916 ABT 119
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a reliable and rapid method to identify differentially expressed genes in microbes. The method relies on the use of a large number of arbitrarily primed PCR reactions. The method is useful for identifying differentially expressed genes in microbes, and for identifying differentially expressed genes in microbes, and for distinguishing genetic differences between two populations of cells which differ in genotype. This method is useful for identifying the DNA sequences of genes involved in the degradation of the picric acid from Rhodooccus exprhropolis strain HL PM-1, and genes involved in cyclobexanol degradation from a consortium of organisms, or to detect cDNA fragments from differentially expressed mRNAs. This method is useful for examining the inhibitory effects of various treaments such as chemicals, environmental pollutants, heavy metals, changes in trate temperature, changes in pH, agents producing oxidative damage, agents producing DNA damage, anaerobiosis, pathogenessis, and changes in nitrate availability on mRNA levels. The present sequence is an universal reamplification primer used in the exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Identifying differentially expressed genes, by amplifying total RNA of first microbial cell population that is contacted with stimulating agent and of a second population using arbitrary primers, and comparing them.
                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Picric acid degradation gene; cyclohexanol degradation; heavy metal; inhibitory effect; chemical; environmental pollutant; anaerobiosis; oxidative damage; pathogenesis; primer; ss.
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   Length 18;
   Score 12.2; DB 1; Length 1
Pred. No. 4.7e+02;
); Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Primer used in the exemplification of the invention.
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ch 2.9%;
1 Similarity 82.4%;
14; Conservative (
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Best Local Similarity
Matches 14; Conserv
   Query Match
Best Local Similarity
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                                                                         Matches
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Nootropic; neuroprotective; antiparkinsonian; anticonvulsant; analgesic; ramquiliser; antidiabetic; ophthalmological; neurodegenerative disorder; neublastin; ischemic neuronal damage; traumatic brain injury; diabetes; peripheral neuropathy; neuropethic pair, Alzheimer's disease; glaucoma; Huntington's disease; Parkinson's disease; amyotrophic lateral sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New truncated neublastin polypeptides lacking one or more amino-terminal amino acids of a mature neublastin polypeptide useful for treating neurodegenerative disorders, e.g. peripheral neuropathy, neuropathic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                            memory impairment; renal disease; PCR; primer; ss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure, Fig 8; 138pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (BIOJ ) BIOGEN INC (NSGE-) NS GENE AS
                                                                                                                                                                                                                                                                                                              WO200272826-A2
                                                                                                                                                                                                                                          Unidentified
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Gaps

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3; Indels

Mismatches

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14; Conservative

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1 AGTCCACGGAGCATATC 17

Human, secreted and transmembrane protein; PRO; virucide; gene therapy; cell death; growth induction cascade; blood coagulation cascade; viral infection; ss.

ted and transmembrane g growth induction cascadion; ss.		-A1.		2001US-00978191.	7US-0062250	7US-0065311	7US-0066364	BUS-0077632	8US-0077641	8US-0077791	BUS-0004022	8US-0078886	8US-0078936	BUS-0078939	8US-0079656	8US-0079664	802-0079689	902-0079786	8US-00/9920	8US-0080105	BUS-0080165	8US-0080194	BUS-0080328	8US-0080333 8US-0080334	98US-0081049P. 98US-0081070P.	8US-0081071	8US-0081203	8US-0081229 8US-0081817	8US-0081819	8US-0081952	8US-0082568	8US-0082569 8US-0082700	8US-0082704	8US-0082804	8US-0082796 8US-0083336	8US-0083322	8US-0083392 8US-0083495	BUS-0083496	8US-0083499	8US-0083545	8US-008355 8US-008355	8US-008355
Human; secre cell death; viral infect	Homo sapiens	US2003050239	13-MAR-2003.	15-OCT-2001;	7-OCT-199	8-NOV-199	1-NOV-199 1-MAR-199	I-MAR-199	I -MAR - 199 I -MAR - 199	2-MAR-199	7-MAR-199	0-MAR-199 0-MAR-199	0-MAR-199	0-MAK-199 5-MAR-199	6-MAR-199	7-MAR-199	7-MAR-199 7-MAR-199	7-MAR-1998	0-MAR-1998	1-MAR-1998	1-MAR-1998	1-MAR-1998 1-APR-1998	1-APR-1998	1-APR-1998 1-APR-1998	08-APR-1998; 08-APR-1998;	8-APR-1998	9-AFR-1998	9-APR-1998 5-APR-1998	5-APR-1998	5-APR-1998	1-APR-1998	1-APR-1998 2-APR-1998	2-APR-1998	2-APR-1998	3-APR-1998 7-APR-1998	8-APR-1998	9-APR-1998 9-APR-1998	9-APR-1998	9-APR-1998 9-APR-1998	9-APR-1998	9-APR-199 9-APR-199	9-APR-199
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98US-0084346P 98US-0084436F 98US-0084411P 98US-0084610P 98US-0084641P 98US-0084641P 98US-0084641P 98US-0084641P 98US-0084641P 98US-0084641P 98US-0084641P 98US-008648P 98US-0085582P 98US-0085582P 98US-008563P 98US-0018441E 98US-0113296F 98US-01131822 98US-01131822 98US-01131822 99US-0123957P 99US-0131822 99US-0131822

30-APR-1998 05-MAY-1998 06-MAY-1998 07-MAY-1998 07-MAY-1998 07-MAY-1998 07-MAY-1998 07-MAY-1998 07-MAY-1998 13-MAY-1998 13-MAY-1998 13-MAY-1998 15-MAY-1998 16-MAY-1998 16-MAY-1998 17-MAY-1998 18-MAY-1998 18-MAY-1998 19-MAY-1998 11-MAY-1998 11-MAY-1998 11-MAY-1998 11-MAY-1998 11-MAY-1998 11-MAY-1998 11-MAY-1998 11-MAY-1999 11-MAY-1998 11-MA

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The specification describes a method for generating genetic disruption (mutation) in bacterial cells. The method comprises providing a bacterial denor cell having a plasmid which comprises a transposable element encoding functions to enable transposition of the transposable element into the host cell nucleic acid and comprising a marker gene and an origin of transpers, and introducing the plasmid from the donor cell to the host cell by conjugation. The method is useful in generating genetic disruptions in bacterial host cells, especially Streptomyces species, and more particularly for generating libraries of bacterial host cells having such disruptions. PCR primers ABZ68345-46 were used to amplify the vph marker from plasmids of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mapping;
                                                                                                                                                                                                                                                                                            Generating a mutation in a bacterial host cell (e.g. Streptomyces spp.), comprises providing a bacterial donor cell (e.g. Escherichia coli) comprising a plasmid and introducing the plasmid to the host cell by conjugation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human, secreted and transmembrane protein, PRO, antiinflammatory, antiarteriosclerotic, cardiant; anti-infertility, anti-HIV, cytostatic, antidiabetic; gene therapy, inflammatory disease; organ failure; atherosclerosis, cardiac injury; infertility; birth defect, premature aging, AIDS, cancer; diabetic complication; chromosome mapping gene mapping; pharmaceitical; diagnostic; biosensor; bioreactor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
2.9%; Score 12.2; DB 1; Length 18;
Best Local Similarity 82.4%; Pred. No. 4.7e+02;
Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 1 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 29; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tissue typing; PCR; primer; ss.
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                                                   24-JJN-2002; 2002WO-GB002884.
                                                                                                 28-JUN-2001; 2001GB-00015894.
                                                                                                                                                  (PLAN-) PLANT BIOSCIENCE LTD
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                                                                                                                                                                                                  Kieser TE;
                                                                                                                                                                                                                                                      WPI; 2003-201505/19.
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09-JAN-2003
                                                                                                                                                                                                     Fowler K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ACA63889;
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Gerritsen ME;
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2000MO-US0346759.
2000MO-US0344956.
2001US-00816744.
2001US-00816220.
2001US-00816220.
2001US-00854208.
                                                                                                                                                                                                                                                2000WO-US005841.
2000WO-US005319.
2000WO-US008439.
2000WO-US013705.
2000WO-US014042.
2000WO-US014941.
2000WO-US015264.
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2000US-00709238.
2000US-00723749.
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Ferrara N, Filvaroff E,
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20-DEC-2000)
20-DEC-2000)
22-MAR-2001)
22-MAR-2001)
22-MAR-2001)
22-MAR-2001)
22-MAR-2001)
25-MAY-2001)
25-MAY-2001)
25-MAY-2001)
26-UN-2001)
26-UN-2001)
29-UN-2001)
29-UN-2001)
29-UN-2001)
29-UN-2001)
                        16-DEC 1999;
30-DEC-1999;
30-DEC-1999;
65-JAN-2000;
66-JAN-2000;
11-FEB-2000;
11-FEB-2000;
11-FEB-2000;
11-MAR-2000;
11-MA
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08-NOV-2000;
27-NOV-2000;
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Matches
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97US-0062250P. 97US-0064249P. 97US-006311P. 97US-0066364P. 98US-0077450P.

03-NOV-1997; 13-NOV-1997; 21-NOV-1997; 10-MAR-1998;

2001US-00999832

24-OCT-2001;

Genetic disruption; mutation; bacterial cell; transposable element; vph;

PCR; primer; ss.

WO2003002738-A1

Synthetic

PCR primer VPH1 used for amplification of vph marker.

22-APR-2003 (first entry)

ABZ68345;

ABZ68345 standard; DNA; 18 BP.

ABZ6834

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17-0CT-1997

19-DEC-2002

US2002192706-A1. Homo sapiens.

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R 11-MAR-1998; 98US-0077441P.

PR 11-MAR-1998; 98US-0077441P.

PR 12-MAR-1998; 98US-0077441P.

PR 12-MAR-1998; 98US-0077491P.

PR 12-MAR-1998; 98US-0077491P.

PR 20-MAR-1998; 98US-007602P.

PR 20-MAR-1998; 98US-007665P.

PR 20-MAR-1998; 98US-007665P.

PR 20-MAR-1998; 98US-007665P.

PR 21-MAR-1998; 98US-0079664P.

PR 27-MAR-1998; 98US-0080137P.

PR 31-MAR-1998; 98US-0080137P.

PR 32-MAR-1998; 98US-0080133P.

PR
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Human; ds; thrombolytic agent; interferon; interleukin; cytokine; erythropoietin; colony stimulating factor; cancer; colorectal carcinoma; apoptosis related condition; AIDS; amyotrophic lateral sollerosis; inflammatory disease; asthma; atherosclerosis; neurodegenerative disease; gastrointestinal disorder; Alzheimer's disease; Parkinson's disease; hypertension; myocardial ischaemia; kidney disease; carcinogenesis; glomerulonephritis; lung disease; pulmonary hypertension; preeclampsia; bronchial asthma; gastric ulcer; renal failure; cardiovascular disease; inflammatory bowel disease; reproductive disorder; premature labour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention describes an isolated nucleic acid (I) comprising, or which is at least 80 % sequence identity to, or the full-length coding sequence of, any of 118 300-2100 nucleotide sequences, which encodes its corresponding PRO polypeptide selected from 18 100-700 amino acid sequences, all given in the specification. The nucleic acids and polypeptides are useful for treating inflammatory diseases, organ failure, atherosclerosis, cardiac injury, infertility, birth defects, premature aging, AIDS, cancer, or diabetic complications. The nucleic acids are useful as hybridisation probes, in chromosome and gene mapping, and in generating antisease RNA or DNA. The polypeptides are useful as pharmaceuticals, diagnostics, biosensors or bioreactors. Both are useful in tissue typing, This sequence represents a novel human secreted and transmembrane PRO polypeptide associated primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                            New secreted and transmembrane nucleic acids and polypeptides, designated as PRO, useful for treating inflammation, organ failure, atherosclerosis, cardiac injury, infertility, birth defects, premature aging, AIDS, or
                                                                                                                                                                                                                                                                                   Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
Goddard A, Godowski PJ, Grimalali JC, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL;
Stewart TA, Tumas D, Williams PM, Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Gaps
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2.9%; Score 12.2; DB 1; Length 18;
Best Local Similarity 82.4%; Pred. No. 4.7e+02;
Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 4 A; 6 C; 5 G; 3 T; 0 U; 0 Other;
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|||||||||||||||||
17 GGAGGTCGACTTCCACT 1
24-AUG-2000, 2000MO-US023328.
01-DEC-2000; 2000MO-US034578.
20-DEC-2000; 2000MO-US034956.
28-FEB-2001; 2001MO-US005520.
22-MAR-2001; 2001MO-US005552.
25-MAY-2001; 2001MO-US017092.
01-JUN-2001; 2001MO-US017980.
20-JUN-2001; 2001MO-US0166.
20-JUN-2001; 2001MO-US01066.
09-JUN-2001; 2001MO-US021066.
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                                                                                                                                                                                                                                             (GETH ) GENENTECH INC.
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98WO-US024855.
98US-00202054.
99WO-US001106.
99US-00254465.
99US-00254465.
99WS-00264213.
99WS-00267213.
99WS-00267213.
99WS-00267213.
99WS-00360133.
99WS-00360133.
                                        97US-0064249P
97US-0064249P
97US-0066341P
98US-0077452P
98US-00776412P
98US-0077641P
98US-0077641P
98US-0077641P
98US-0077641P
98US-0077641P
98US-0077641P
98US-0077641P
98US-0077641P
98US-0077641P
98US-0077664P
98US-0079664P
                                                                                                                                                                                                                                                                                                                                     2000WO-US003565.
2000WO-US004341.
2000WO-US005004.
                                                                                                                                                                                                                                                                                                                          2000WO-US000277
2000WO-US000376
                               15-OCT-2001; 2001US-00978192
          US2002177553-A1.
Homo sapiens.
                     28-NOV-2002,
                                                                                                                                                                                                  20-NOV-1998
                                                                                                                                                                                  07-OCT-19
02-NOV-19
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The invention relates to an isolated secreted and transmembrane polypeptide, designated as PRO polypeptide is useful for in PRO polypeptide is useful for in PRO polypeptide or an including a bioactive molecule to a cell. The PRO polypeptide or an atibody against it is useful for modulating a biological activity of a cell. The PRO polypeptide is useful for modulating a biological activity of a natibody against it is useful for modulating a biological activity of a cell. The PRO polypeptide is useful for modulating factor and other polypeptide is also useful as a thrombolytic agent, interferon, interferon, onlypeptide is useful for treating disease such as the PRO polypeptide is useful for treating disease such as cancer e.g. colorectal carcinoma; apoptosis related conditions e.g. AIDS, atherosclerosis, neurodegenerative disease e.g. Alzheimer's disease, atherosclerosis; neurodegenerative disease e.g. Alzheimer's disease, atherosclerosis; neurodegenerative disease e.g. Alzheimer's disease, comparation is chaemai, kidney disease e.g. pulmonary hypertension, bronchial achieval and disease, reproductive disease e.g. pulmonary hypertension, bronchial asthem associated olisorders e.g. pulmonary hypertension, bronchial achieval associated olisorders e.g. premature labour and preeclampsia, carcinogenesis. The present sequence represents a PRO polypeptide associated disease. G. premature labour and preeclampsia, carcinogenesis. The present sequence represents a PRO polypeptide associated disease. The present disease e.g. the printed sequence data for this patent did not form part of the printed sequence thim PDOCID=20020177553
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated PRO polypeptides e.g. PRO213, PRO274 and PRO300, for use as pharmaceuticals, diagnostics, biosensors and bioreactors, for identifying modulators of receptor-ligand interactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL;
Stewart TA, Tumas D, Williams PM, Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; SEQ ID NO 519; 55pp; English.
                                                            27-NOY-2000; 2000US-00123749.
01-DEC-2000; 2000WG-US034278.
20-DEC-2000; 2000WG-US034256.
22-MAR-2001; 2001WG-US034256.
22-MAR-2001; 2001WS-00815744.
22-MAR-2001; 2001WS-00815744.
22-MAR-2001; 2001WS-00815744.
22-MAR-2001; 2001WS-00815780.
10-MAY-2001; 2001WS-00815780.
10-WAY-2001; 2001WS-00815780.
10-UNN-2001; 2001WS-0081581.
10-UNN-2001; 2001WS-0081581.
11-UNN-2001; 2001WS-0081581.
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Gaps .; 0

Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels

149 GGAGGCCGGCTTCGACT 165

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GGAGGTCGACTTCCACT 1

17

Sequence 18 BP; 4 A; 6 C; 5 G; 3 T; 0 U; 0 Other;

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Page 442

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Human, PRO polypeptide, secreted and transmembrane protein, immune disorder, diabetes, hyper-insulinaemia, hypo-insulinaemia, cardiac insulficiency, nervous system disorder, kidney disorder; bone disorder, carthiage disorder, arthritis; tumour, wound healing; genetic disorder; cytostatic, antidiabetic; antinflammatory, antiarthritic, anti-tumour, vulnerary, antianaemic, dermatological; cardiant; PCR; primer; ss.
                                                                         Human PRO DNA PCR primer SEQ ID No 519
                                                                                                                                                                                                                                          81US-00267213.
97US-0062280P.
97US-0064249P.
97US-0065311P.
97US-0074250P.
98US-0077449P.
98US-0077449P.
98US-00777449P.
98US-00777449P.
98US-00777449P.
98US-00777449P.
98US-0078924P.
98US-0079689P.
98US-0079689P.
98US-0079689P.
98US-0079689P.
98US-0079644P.
98US-0079689P.
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98US-0079689P.
98US-0079688P.
98US-0079689P.
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98US-0079689P.
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98US-0079688P.
98US-0079688P.
98US-0079688P.
                   ABX92693 standard; DNA; 18 BP
                                                                                                                                                                                                                          2001US-00978697
                                                        entry)
                                                       (first
                                                                                                                                                                                     US2002169284-A1.
                                                                                                                                                                    Homo sapiens.
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17-OCT-1997;
03-NOV-1997;
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11. MAR. 1998;
11. MAR. 1998;
12. MAR. 1998;
13. WAR. 1998;
17. MAR. 1998;
20. MAR. 1998;
20. MAR. 1998;
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25-MAR-1998;
26-MAR-1998;
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05-MAR-1999;
08-MAR-1999;
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27-MAR-1998;
27-MAR-1998;
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27-MAR-1998;
30-MAR-1998;
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02-NOV-1998;
06-NOV-1998;
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26-JUN-1998;
07-OCT-1998;
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                                                      08-MAY-2003
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                                     ABX92693;
RESULT 865
ABX92693/c
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I, Baker KP, Botstein D, Desnoyers L, Eaton D; Filvaroff E, Fong S, Gao W, Gerber H, Gerriteen ME; Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ; Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton Tumas D, Williams PM, Wood WI;
                                                              2000WO-US000277

2000WO-US000376

2000WO-US000376

2000WO-US005841

2000WO-US005841

2000WO-US005841

2000WO-US005843

2000WO-US00532

2000WO-US014941

2000WO-US014941

2000WO-US012564

2000WO-US012564

2000WO-US012564

2000WO-US012564

2000WO-US012564

2000WO-US01264

2000WO-US01264

2000WO-US01264

2000WO-US02652

2000WO-US03865

2000WO-US03865

2000WO-US03865

2000WO-US03865

2001WO-US03865

2001WO-US03865

2001WO-US03865

2001WO-US03865

2001WO-US03865

2001WO-US03865

2001WO-US086520

2001WO-US086520

2001WO-US086520

2001WO-US086520
                                                                                                                                                                                                                                                                                                                       2001NO-US017092.
2001US-00872035.
2001NO-US017800.
2001US-00874503.
2001US-00886342.
         99WO-US028551.
99WO-US028565.
99WO-US030095.
99WO-US031243.
                                                        2000WO-US000219.
                                                                                                                                                                                                                                                                                                                                                                                                    09-JUL-2001; 2001WO-US021735
30-JUL-2001; 2001US-00918585
                                                                                                                                                                                                                                                                                                                                                                                                                                  (GETH ) GENENTECH INC.
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Ferrara N, Fi
Goddard A, Go
Kljavin IJ, K
Stewart TA, T
        02-DEC-1999;
02-DBC-1999;
16-DBC-1999;
30-DBC-1999;
30-DBC-1999;
06-JAN-2000;
06-JAN-2000;
06-JAN-2000;
06-JAN-2000;
11-FEB-2000;
24-FEB-2000;
10-MAR-2000;
31-MAR-2000;
31-MAR-2000;
22-MAY-2000;
22-MAY-2000;
                                                                                                                                                                        30-MAY-2000;
02-JUN-2000;
28-JUL-2000;
24-AUG-2000;
08-NOV-2000;
27-NOV-2000;
                                                                                                                                                                                                                                01-DEC-2000;
20-DEC-2000;
20-DEC-2000;
28-FEB-2001;
22-MAR-2001;
22-MAR-2001;
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10-MAY-2001;
10-MAY-2001;
25-MAY-2001;
01-JUN-2001;
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05-JUN-2001;
14-JUN-2001;
19-JUN-2001;
20-JUN-2001;
29-JUN-2001;
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<u>D</u>I;

Novel secreted and transmembrane polypeptides and polynucleotides encoding them useful for treating cancer, kidney diseases, bone, cartilage disorders and immune deficiencies.

Example 95; Page 179; 459pp; English.

The present invention relates to the isolation of novel human PRO polypeptides, and the polynucleotide sequences encoding them. The PRO polypeptides are secreted and transmembrane proteins. The PRO polypeptides are useful for detecting other PRO polypeptides, for linking bioactive molecules to calls expressing PRO polypeptides, for modulating biological activities of calls expressing PRO polypeptides, and for for identifying agonists or antagonists. The bioactive molecule maybe a toxin, radiolabel or antibody, and causes apoptosis or death of the cell. The PRO polypeptides are useful for treating immune disorders, diabetes of hyper. Insulinaemia, cardiac insufficiency, narrous system disorders, kidney disorders, bone and cartilage disorders or arthritis, tumours, and wound healing. The polynucleotide sequences encoding PRO polypeptides are useful as hybridisation probes, in chromosome and gene

99US-00380137. 99US-00380138. 99US-00380142.

25-AUG-1999; 25-AUG-1999; 25-AUG-1999;

99US-00284291

10-MAR-1999; 10-MAR-1999; 12-APR-1999;

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mapping, in the generation of antisense RNA and DNA, in the preparation of PRO polypeptides, for generating transgenic animals or knockout animals, for the genetic analysis of individuals with genetic disorders, and in gene therapy. The present sequence represents a PCR primer used in the examples of the present invention. Note: The sequence data for this patent was obtained in electronic format directly from the USPTO web site at sequence.
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Sequence 18 BP; 4 A; 6 C; 5 G; 3 T; 0 U; 0 Other;

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0
2.9%; Score 12.2; DB 1; Length 18; 82.4%; Pred. No. 4.7e+02; tive 0; Mismatches 3; Indels
                                                      149 GGAGGCCGGCTTCGACT 165
                                                                      17 GGAGGTCGACTTCCACT
                           Conservative
                Similarity
                             14;
  Query Match
Best Local S
                 Best Loc
Matches
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Human HBM STS marker forward primer #118. BP. ACC45656 standard; DNA; 18 (first entry) 02-JUN-2003 ACC45656; 998 RESULT 86 ACC45656/

Human, high bone mass, HBM, LRP5, LRP6, transgenic; bone mass modulation, gene therapy; bone density modulation, bone size, some clistue connectivity, bone disease; osteoporosis, PCR, osteomalacia, rickets, Paget's disease; neoplasm of the bone; primer; ss.

Homo sapiens.

WO200292764-A2 21-NOV-2002.

11-MAY-2001; 2001US-0290071P. 13-MAY-2002; 2002WO-US014876

17-MAY-2001; 2001US-0291311P. 01-FEB-2002; 2002US-0353058P. 04-MAR-2002; 2002US-0361293P

Bodine PV; (GENO-) GENOME THERAPEUTICS CORP. Yaworsky PJ, Bex FJ, AMHP) WYETH. Babij P,

WPI; 2003-129278/12.

New transgenic animals (e.g. mice), useful as models for studying bone density modulation, developing drugs for treating or preventing bone diseases (e.g. osteoporosis), or diagnosing diseases characterized by reduced bone density.

Disclosure; Page 55; 603pp; English.

The invention relates to novel transgenic animals expressing the high bone mass (HBM) gene, expressing the corresponding wild type HBM gene, comprising an alteration of the gene encoding inFS or LRP6, or expressing an IMPE that is modulated by an altered gene control sequence introduced by homologous or non-homologous recombination. The transgenic animals are for the study of bone density modulation or bone mass modulation. The invention has osteopathic and cytostatic activity. The polynucleotides of the invention may have a use in gene therapy. The transgenic animals and nucleic acids are for the study of bone density modulation, where the bone mass is modulated relative to non-transgenic animals of the same species in more than one parameter selected from bone density, bone strength, trabecular number, bone size, or bone tissue connectivity. The

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transgenic animals, nucleic acids and methods are useful for identifying molecules involved in bone development, and for developing pharmaceutical compositions, which may be employed for treating or preventing bone diseases. e.g. osteoporosis, osteomalacia, rickets, Paget's disease, or neoplasms of the bone. The transgenic animals and nucleic acids are also useful in methods for diagnosing diseases involved in bone development, or characterised by reduced bone density or mass. The present sequence is
                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                        ;
0
                                                                                                                                                                                                                                                    Match 2.9%; Score 12.2; DB 1; Length 18; Local Similarity 82.4%; Pred. No. 4.7e+02; Les 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                               Sequence 18 BP; 3 A; 8 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                   used in the exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                  18 GGGCAGGAGTGACTCTG
                                                                                                                                                                                                                                                                                                                                                       1 GGCCAGGAGTGAAACTG
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                                                                                                                                                                                                                                                                                                             Matches
       8888888888
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Human, mouse, 88; probe; gene 216; antiasthmatic; antiinflammatory; anorectic; chromosome 20p13-p12; single nucleotide polymorphism; SNP; gene therapy; respiratory disease; asthma; obesity; bronchial hyper-responsiveness; chronic obstructive pulmonary disease; adult respiratory distress syndrome; inflammatory bowel syndrome. Human 216 gene allele specific oligonucleotide probe #39. WO200283077-A2. Homo sapiens. 24-OCT-2002

В

ABX75208 standard; DNA; 18

RESULT 867

(first entry)

25-MAR-2003

ABX75208;

15-APR-2002; 2002WO-US012063. 13-APR-2001; 2001US-00834597. 13-APR-2001; 2001WO-US012245.

(SCHE) SCHERING CORP. (GENO-) GENOME THERAPEUTICS CORP.

Del Mastro RG; Dupuis J, Van Eerdewegh P, Pandit S; Little RD, Allen K, F Keith T, Simon J,

WPI; 2003-092960/08

ö New isolated gene 216 nucleic acids, useful for diagnosing, preventing treating a disorder, such as asthma, bronchial hyper-responsiveness, chronic obstructive pulmonary disease, obesity or inflammatory bowel syndrome.

Example 10; Page 166; 650pp; English.

This invention relates to a novel isolated nucleic acid, gene 216, identified from human chromosome 20p13-pi2. The invention also discloses regions of the 216 gene that contain single nucleotide polymorphisms (SNP's) which may be used as markers for disease susceptibility or severity. The nucleotides of the invention may have antiasthmatic, antiminammatory or anorectic activities and may be used in gene therapy. The nucleic acids, antibodies or its fragments are useful for diagnosing, preventing or treating a disorder, such as respiratory diseases (e.g. asthma, bronchial hyper-responsiveness, chronic obstructive pulmonary disease or adult respiratory distress syndrome), obesity, or inflammatory bowel syndrome. The nucleic acids are also useful for identifying

increased susceptibility of a subject to the disorders mentioned. The nucleic acids can also be used as primers and templates for the recombinant production of disorder-associated peptides or polypeptides, for chromosome and gene mapping, or for tissue distribution studies. The present sequence represents a gene 216 specific oligonucleotide probe used in the scope of the invention 88888888888

Sequence 18 BP; 1 A; 3 C; 9 G; 5 T; 0 U; 0 Other;

Gaps ö 2.9%; Score 12.2; DB 1; Length 18; 82.4%; Pred. No. 4.7e+02; ative 0; Mismatches 3; Indels Best Local Similarity 82.4 Matches 14; Conservative Query Match

ö

92 CATCACCACGICIGACC 108

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17 CAGCACCACAGCTGACC

RESULT 868 ABZ79946/

ABZ79946 standard; DNA; 18

BP.

ABZ79946;

19-MAY-2003 (first entry)

Mycobacterium tuberculosis rpsI PCR primer SEQ ID NO:16.

Mycobacterium tuberculosis, mutT2; alkh; ogt; Rv3908; mutY; Rv3909; detection; multidrug resistance; multiple drug resistance; MDR; infection; PCR primer; ss.

Mycobacterium tuberculosis.

Synthetic

WO2003016562-A2

27-FEB-2003

14-AUG-2002; 2002WO-EP009679,

14-AUG-2001; 2001US-0311824P. 21-AUG-2001; 2001US-0313523P.

(INSP) INST PASTEUR

Gicquel B;

WPI; 2003-256711/25

Predicting the epidemic character of a Mycobacterium tuberculosis isolate and/or the acquisition of multiple drug resistance (MDR) by the isolate by detecting an alteration in the DNA repair system of the isolate.

Disclosure; Page 17; 83pp; English.

The present invention describes a method for predicting the epidemic character of a Mycobacterium tuberculosis isolate and/or a selective advantage to be maintained in the host and/or the acquisition of multiple drug resistance (MDR) by the isolate comprising detecting an alteration in the DNA repair system of the isolate. Also described: (1) detecting a pythomorphy (12) a polymucleotide; (3) a kit for detecting Mycobacterium tuberculosis; (4) an Escherichia coli strain containing the plasmid pMYC2501; and (5) an Escherichia in a patient infected by Mycobacterium tuberculosis a higher risk of being unable to eliminate the bacillus or of developing MDR tuberculosis. The method is useful for predicting the epidemic character of a Mycobacterium tuberculosis isolate and/or a selective advantage to be maintained in the host and/or the acquisition of MDR by the isolate. The present sequence represents a PCR primer for M. tuberculosis rpsI, which is used in the exemplification of the present invention

BP; 4 A; 5 C; 5 G; 4 T; 0 U; 0 Other; Sequence 18

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human, 88; PCR; secreted protein; transmembrane protein; PRO; primer; malignancy; cancer; ovarian cancer; colorectal cancer; sarcoma; leukaemia; lymphoma; inflammatory disease; necrosis; atherosclerosis; infertility; premature aging; psoriasis; inflammatory disease; renal disease; arthritis; immune-mediated alopecia; stroke; encephalitis; hepatitis; multiple sclerosis; gene therapy.
                                                  Gaps
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0
Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                      Human secreted/transmembrane protein PRO298 PCR primer #3.
                                                                                                 GGCACCAAGCTGGTGAA 298
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97US-0064341P
97US-0066341P
98US-0077450P
98US-0077641P
98US-0077641P
98US-0077649P
98US-0077649P
98US-0077649P
98US-007763P
98US-0078930P
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98US-00184216.
98US-00187368.
98WO-US024855.
98US-00202054.
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ACA66434 standard; DNA; 18
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11.-MAR-1998;
12.-MAR-1998;
13.-MAR-1998;
17.-MAR-1998;
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26-JUN-1998;
07-OCT-1998;
07-OCT-1998;
02-NOV-1998;
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2001US-00854280.
2001WO-US017092.
2001US-00872035.
2001WS-00874503.
2001US-00884503.
    9905-00311832-9900-00311832-9900-00310733-9905-00380138-9905-00380138-9900-00302851-9900-003031243-9900-0031243-
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2001US-00816920.
2001WO-US009552.
2001US-00854208.
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2000WO-US013705.
2000WO-US014042.
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2000WO-US032678.
2000US-00747259.
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2000WO-US003565.
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2000WO-US006319.
2000WO-US007532.
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2000WO-US030873.
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2001WO-US006520
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25-MAY-2001; 200
01-JUN-2001; 200
                                                                                 05-JAN-2000;
06-JAN-2000;
06-JAN-2000;
11-FEB-2000;
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10-MAR-2000;
21-MAR-2000;
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17-MAY-2000;
22-MAY-2000;
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02-JUN-2000;
28-JUL-2000;
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01-DEC-2000;
20-DEC-2000;
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08-NOV-2000;
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GETH) GENENTECH INC.

Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL; Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME; Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ; Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL; Stewart TA, Tumas D, Williams PM, Wood WI;

WPI; 2003-341189/32.

New genes and secreted and transmembrane polypeptides (e.g. PRO337 or PRO1559), useful for treating or diagnosing e.g. cancers, atherosclerosis, infertility, stroke, encephalitis, hepatitis or multiple sclerosis in mammals.

Example 95; Page 180; 460pp; English.

The invention relates to a new isolated nucleic acid molecule comprises a sequence with at least 80% identity to: (a) a nucleotide encoding any of 94 PRO polypeptides whose sequences are fully defined in the

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cc specification; or (b) any of 94 mucleotide sequences fully defined in the specification; or the full length coding sequence of any these 94 concleted sequences. Also included are an isolated PRO polypeptide concletions sequences an isolated PRO polypeptide having at least 80% paintives when compared to any of the PRO polypeptide having cf 1 least 80% paintives when compared to any of the PRO polypeptide having cf 2 at least 80% anno acid sequence identity to: (a) an amino acid sequence concleted with ATCC numbers listed in the specification; (b) the PRO polypeptide, lacking its associated signal peptide, a colon acid sequence and any of the PRO polypeptide, with or comprising its associated signal peptide; a colonaria period of comprising the PRO polypeptide, with or colorecule, a host cell comprising the vector comprising the nucleic acid molecule, a host cell comprising the vector (and producing a PRO colorecule, a chieracle sequence and a nati-PRO antibody. The PRO polypeptides or polymucleotides are useful as pharmaceuticals, colorecule or polymucleotides are useful as pharmaceuticals, colorectal cancer, sarcoma, leukaemia or lymphoma; inflammatory disease, colorectal cancer, sarcoma, infertility, premature aging, postiasis, colorectal cancer, sarcoma, infertility, premature aging, postiasis, inflammatory disease, colorectal cancer, sarcoma, infertility, premature aging, postiasis, colorectal cancer, sarcoma, infertility, premature aging, postiasis, inflammatory disease, renal disease, renal disease, renal disease, colorectal cancer, sarcoma, infertility, premature aging, postiasis, colorectal cancer, sarcoma, independent and presence of these diseases; The PRO polypeptides are determination of the prosence acidentical cancer, sarcom
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97US-0064249P.
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98US-00777649P.
98US-0077791P.
98US-00788186P.
98US-0078819P.
98US-0078819P.
98US-0078634P.
98US-0079664P.
98US-0079664P.
98US-0079664P.
98US-0079728P.
98US-0079728P.
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98US-0081337P.
98US-0081347P.
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98US-0081339P.
98US-0084649P.
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33-MAR-1998;
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98US-0086744P.
98US-0086744P.
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98US-0086414P.
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98US-0191326P.
98US-0191326P.
98US-0191325P.
98US-0

15-MAY-1998;
15-MAY-1998;
22-MAY-1998;
22-MAY-1998;
22-MAY-1998;
22-MAY-1998;
23-MAY-1998;
24-MAY-1998;
25-MAY-1998;
26-UJM-1998;
26-UJM-1998;
27-MAY-1998;
26-UJM-1998;
27-MAY-1998;
28-MAY-1998;
28-MAY-1998;
28-MAY-1999;
30-NOV-1999;
30-NO

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Ashkenazi (GETH)

GENENTECH INC Baker

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Filvaroff E, Fong S, Gao W, Gerber H, Garritsen ME, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ; Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL; Tumas D, Williams PM, Wood WI,
                                                                                                                    New isolated PRO polypeptides for example extracellular, secreted and membrane bound proteins, useful for modulating the biological activities of cells and for treating, for example diabetes, cancer, rheumatoid arthritis, and hearing loss:
                                                                                                                                                                                                                                  The invention describes an isolated secreted and transmembrane (PRO) polypeptide (I). PRO337 polypeptide is useful for detecting PRO4993 useful for detecting PRO4993 useful for detecting PRO155 polypeptide in a sample, and PRO155 are useful for detecting PRO155 polypeptide in a sample, and PRO1559 is useful for detecting PRO355, PRO700 and PRO39 in a sample, and PRO393 is useful for linking a bioactive molecule to a cell expressing a PRO337 polypeptide, and PRO337 is useful for linking a bioactive molecule concludentle to a cell expressing a PRO4993 polypeptide. PRO1559 is useful for linking a bioactive molecule to a cell expressing a PRO493 polypeptide. PRO7155 PRO700 and PRO739 polypeptides, and PRO735, PRO700 and PRO739
                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                       ch 2.9%; Score 12.2; DB 1; Length 18; 1 Similarity 82.4%; Pred. No. 4.7e+02; 14; Conservative 0; Mismatches 3; Indels
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97US-006424PP.
97US-006634LP.
97US-0066364P.
98US-0077632P.
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98US-0077641P.
98US-0078044P.
98US-0078044P.
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                                                                                     WPI; 2003-521814/49.
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   Ferrara N, F
Goddard A, G
Kljavin IJ,
Stewart TA,
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13-MAR-1998;
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20-MAR-1998;
20-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-NOV-1997;
13-NOV-1997;
21-NOV-1997;
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Best Local
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ACD30035/c
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PR 20-MRR-1998 PUS-0079639P.
PR 25-MRR-1998 PUS-0079663P.
PR 27-MRR-1998 PUS-0079728P.
PR 21-MRR-1998 PUS-0079728P.
PR 31-MRR-1998 PUS-0079728P.
PR 31-MRR-1998 PUS-007973P.
PR 31-MRR-1998 PUS-007973P.
PR 31-MRR-1998 PUS-007973P.
PR 31-MRR-1998 PUS-0080117P.
PR 01-APR-1998 PUS-008117P.
PR 01-APR-1998 PUS-008117P.
PR 01-APR-1998 PUS-008179F.
PUS-0081
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The invention describes an isolated, secreted and transmembrane polypeptide, termed PRO polypeptide (1). (1) is useful for detecting throughly PROJ37, PROJ37, PROJ37, PROJ37, PROJ37, PROJ37, PROJ37, PROJ37, PROJ37, and for linking a bioactive molecule to a cell expressing the above polypeptides. The bloactive molecule is a toxin, radiolabel or an antibody and causes cell death. (1) is useful as therapeutic agent, in medical and industrial applications e.g. for treating neuropathy, especially peripheral neuropathy, disbetic peripheral neuropathy, AIDS-associated neuropathy, Charcot-Marie-Tooth disease, Refusum's disease, Abetalipoproteinaemia, Tangier disease, Krabbe's disease, Metachromatic leukodystrophy, Fabry's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer;
                           Novel secreted and transmembrane polypeptide for modulating biological activity of cell expressing the polypeptide, identifying agonists or antagonists of polypeptide, and as molecular weight markers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer; ss; inflammatory disease; organ failure; atherosclerosis; cardiac injury; infertility; birth defect; premature aging; AIDS; diabetic complication; tissue typing; human; PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human secreted/transmembrane polypeptide PRO298 primer #3
                                                                                                                         Example 95; Page 177; 459pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                149 GGAGGCCGGCTTCGACT 165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GGAGGTCGACTTCCACT 1
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22-MAY-1998; 98US-0086486P-
28-MAY-1998; 98US-0087098P-
28-MAY-1998; 98US-0087098P-
28-MAY-1998; 98US-0087098P-
26-UUN 1998; 98US-0087208P-
31-UUL-1998; 98US-0090863P-
26-UUN 1998; 98US-0090863P-
26-UUN 1998; 98US-010908P-
31-SEP-1998; 98US-010908P-
22-NOV-1998; 98US-0109304P-
22-NOV-1998; 98US-0109304P-
22-NOV-1998; 98US-0109304P-
23-NOV-1998; 98US-010320P-
23-DEC-1998; 98US-010320P-
23-DEC-1998; 98US-010320P-
23-DEC-1998; 98US-013022P-
23-DEC-1998; 98US-013022P-
23-DEC-1998; 98US-013022P-
23-DEC-1999; 99US-013022P-
23-DEC-1999; 99US-013022P-
24-MAY-1999; 99US-0131022P-
25-DEC-1999; 99US-013022P-
26-DEC-1999; 99US-0131022P-
26-DEC-1999; 99US-01310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GETH ) GENENTECH INC.
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WPI; 2003-503575/47.

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PR 30-MAR-1998 98US-00090105P
PR 31-MAR-1998 98US-00801105P
PR 31-MAR-1998 98US-00801105P
PR 11-MAR-1998 98US-00801105P
PR 10-APR-1998 98US-00801105P
PR 10-APR-1998 98US-0080134P
PR 10-APR-1998 98US-0081137P
PR 10-APR-1998 98US-008119P
PR 20-APR-1998 98US-008119P
PR 20-APR-1998 98US-008119P
PR 20-APR-1998 98US-008119P
PR 20-APR-1998 98US-0081342P
PR 20-APR-1998 98US-008133P
PR 20-APR-1998 98US-008139P
PR 20-APR-19
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PR 07-OCT 1998; 98105-00169378.

PR 07-OCT 1998; 9880-00183168.

PR 06-NOV-1998; 98105-00183168.

PR 20-NOV-1998; 98105-00183128.

PR 20-NOV-1998; 98105-0018328.

PR 20-NOV-1998; 98105-0013821.

PR 20-NOV-1998; 98105-0013828.

PR 20-NOV-1998; 98105-0013828.

PR 20-NOV-1998; 98105-0018328.

PR 20-NOV-1999; 98105-0026513.

PR 20-NOV-1999; 98105-0026513.

PR 20-NOV-1999; 98105-0026513.

PR 20-NOV-1999; 98105-0026513.

PR 20-NOV-1999; 99105-0026513.

PR 20-NOV-1999; 99105-0126773.

PR 20-NOV-1999; 99105-013622.

PR 20-NOV-1999; 99105-013622.

PR 20-NOV-1999; 99105-013622.

PR 20-NOV-1999; 99105-013622.

PR 20-NOV-1999; 99105-013613.

PR 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; secreted and transmembrane protein; PRO; viral infection; trumour growth; retinal disorder; injury; sight loss; retinal disorder; injury; sight loss; retinal growth; retinal disorder; injury; sight loss; sport-related joint problem; age-related macular degeneration; sport-related joint problem; articular cartilage defect; osteoarthritis; rheumarcioid arthritis; wound healing; obselty; diabetes; insulinaemia; kidney disorder; mesangial cell function; Berger disease; nephropathy; cellac disease; dermatitis; Crohn disease; neuropathy; diabetic peripheral neuropathy; peripheral neuropathy; reduced motility of the gastrointestinal tract; reduced motility of the gastrointestinal tract; charled the urinary bladder; post polio syndrome; Krabbe's disease; Refsum's disease; Pabry's disease; Tangier disease; Refsum's disease; Tangier disease;
                                                                                                                                                                  Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL,
Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel human secreted and transmembrane protein related primer #220
                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                              Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 873
ACD29450 standard; DNA; 18 BP.
XX
ACD29450;
XX
T 27-AUG-2003 (first entry)
XX
Human; secreted and transmembrant
XW
Tumour growth; retinal disorder;
XW
Tetinitis pigmentosum; age-relate
XW
Totalac disease; darmatis; croh
XW
Totalac insulficiency disorder; with a cardiac motility of the gastroin
XW
Totalac motility of the gastroin
XW
Totalac motility of the gastroin
XW
Charcot-Maria-Tooth disease; Fab;
XX
XX
Homo sapiens.
XX
D3-MAR-2003.
XX
TO-CT-1997; 97US-0065250P.
PR
13-MAR-2003.
XX
TO-CT-1997; 97US-006534P.
PR
13-NOV-1997; 97US-006534P.
PR
13-NOV-1997; 97US-006534P.
PR
13-NOV-1997; 97US-00634P.
PR
13-NOV-1997; 97US-00634P.
PR
13-NOV-1997; 97US-00634P.
PR
13-NAR-1998; 98US-0077641P.
PR
13-MAR-1998; 98US-0077641P.
PR
20-MAR-1998; 98US-0077641P.
PR
21-MAR-1998; 98US-0077641P.
PR
21-MAR-
                                                                                                                                                                                                                                                                                                                149 GGAGGCCGGCTTCGACT 165
19-JUN-2001; 2001US-00886342.
20-JUN-2001; 2001WO-US019692.
29-JUN-2001; 2001WO-US021735.
30-JUL-2001; 2001US-00918585.
                                                                                                                                                                                                                                                                                                                                         17 GGAGGTCGACTTCCACT 1
                                                                                                                          (GETH ) GENENTECH INC
    qq
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9908-0130232P.
9908-0131022P.
9908-01311428P.
9908-01314287P.
9908-01314287P.
9908-01313557P.
9908-0139557P.
9908-0140580P.
9908-0140580P. 9805-0094651P9805-0100038P9805-010842169805-001842169805-001873689805-001873689805-001851P9805-0113621P9805-0113621P9905-0136289906-0136289906-0136289906-0136289906-0136289906-0136289906-0136289906-0136289906-0136289906-0136289906-0136289906-013628-99US-0123957P. 99US-0126773P. 99US-00284291. 10-MAY-2001; 2001US-00854280. 25-MAY-2001; 2001WO-US017092. 01-JUN-2001; 2001US-00872035. 2000WO-US000376 2000WO-US004341 2000WO-US005004 2000WO-US000277 98US-0091359P 30-MAY-2000; 02-JUN-2000; 28-JUL-2000; 24-AUG-2000; 08-NOV-2000; 27-NOV-2000; 11-FEB-2000; 18-FEB-2000; 24-FEB-2000; 25-AUG-1999; 25-AUG-1999; 25-AUG-1999; 29-OCT-1999; 30-NOV-1999; 02-DEC-1999; 07-0CT-1998 07-0CT-1998 02-NOV-1998 06-NOV-1998 20-NOV-1998 07-DEC-1998 22-DEC-1998 23-DEC-1998 23-DEC-1998 05-JAN-19 05-MAR-19 08-MAR-19 02-JUN-

The present invention relates to the isolation of a human tummour suppressor gene, TSLL1 (hTSLL1), and the encoding protein. The TSLL1 gene and protein are useful for preventing and treating cancers. The gene is useful for diagnosing carcinoma in pre-critical stages, qualitative diagnosis of carcinoma, predicting the prognosis of cancer therapy, and forecasting the sensitivity of a carcinoma to chemical therapy, and radiotherapy and gene therapy. The TSLL1 protein is homologous the TSLC1 protein. The present sequence represents a PCR primer used to generate a probe for human TSLL1 CDNA. New protein encoded by tumor suppressor gene, designated as TSSL1 gene, useful for preventing or treating cancers, predicting of prognosis of cancer therapy, or diagnosing carcinoma in pre-clinical stages. Human tumour suppressor gene; TSLL1; hTSLL1; cancer; carcinoma; pre-critical stage; cancer therapy; chemical therapy; radiotherapy; TSLC1; PCR; primer; 88. Sequence 18 BP; 5 A; 7 C; 4 G; 2 T; 0 U; 0 Other; (PRES-) PRESIDENT NAT CANCER CENT. (BMLB-) BML INC. Example; Page 6; 20pp; English. 29-AUG-2002; 2002US-00230335. 11-OCT-2001; 2001JP-00313966. Murakami Y, Nomura S; WPI; 2003-626209/59. US2003109016-A1. Homo sapiens 12-JUN-2003.

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Gaps

.. 0

Length 18; 3; Indels

Score 12.2; DB 1; Pred. No. 4.7e+02; 0; Mismatches 3;

Query Match 2.9%; Best Local Similarity 82.4%; Matches 14; Conservative

01-JUN-2001; 2001WO-US017800. 05-JUN-2001; 2001US-00874503. 14-JUN-2001; 2001US-00882636.

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149 GGAGGCCGGCTTCGACT 165

ઠ g

17 GGAGGICGACTICCACT 1

PCR primer #1 for generating human TSLL1 probe.

20-NOV-2003 (first entry)

ADA24424;

ADA24424 standard; DNA; 18 BP.

RESULT 874

Arcrecaederecire 2 18

RESULT 875 ADB98354/C

334 ACGACCAGGCCGGCTG 350

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Gaps

6

2.9%; Score 12.2; DB 1; Length 18; 82.4%; Pred. No. 4.7e+02; ative 0; Mismatches 3; Indels

Query Match Best Local Similarity 82.4 Matches 14; Conservative

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antibacterial; immunosuppressive; neuroprotective; PCR; primer; ss
                                                                                                                                            9705-0064249P.
9705-0064249P.
9705-0065644P.
9705-0065644P.
9705-0077450P.
9805-0077641P.
9805-0077641P.
9805-0077641P.
9805-0077641P.
9805-0077641P.
9805-0079294P.
9805-0080327P.
9805-0080327P.
9805-0080327P.
9805-0080334P.
9805-0080334P.
9805-0081238P.
9805-0081338P.
9805-0081338P.
9805-0081338P.
9805-0081338P.
9805-0081338P.
9805-0081338P.
9805-0081338P.
9805-008338P.
9805-008338P.
9805-008338P.
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98US-0083742P.
98US-0084366P.
                                                                                                                   16-OCT-2001; 2001US-00978608
                                                          US2003045462-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-APR-1998;
30-APR-1998;
05-MAY-1998;
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22-APR-1998;
22-APR-1998;
22-APR-1998;
23-APR-1998;
27-APR-1998;
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11 - MAR - 1998

11 - MAR - 1998

11 - MAR - 1998

12 - MAR - 1998

13 - MAR - 1998

14 - MAR - 1998

20 - MAR - 1998

20 - MAR - 1998

25 - MAR - 1998

25 - MAR - 1998

27 - MAR - 1998
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31-MAR-1998;
31-MAR-1998;
31-MAR-1998;
01-APR-1998;
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01-APR-1998;
08-APR-1998;
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09-APR-1998;
15-APR-1998;
15-APR-1998;
15-APR-1998;
15-APR-1998;
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30-MAR-1998;
30-MAR-1998;
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21-APR-1998;
21-APR-1998;
                               Homo sapiens
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                                                                                        06-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to High Bone Mass (HBM), LRP5 (Zmax1) and LRP6 mutants, which results in a HBM-like phenotype when expressed in a cell. The HBM-like phenotype results in bone mass modulation and/or lipid level modulation. The invention is useful for diagnosing a HBM-like phenotype in a subject and for preparing a composition for modulating bene mass and/or lipid levels in a subject suffering from e.g. osteoporosis. The present sequence is a Sequence Tagged Site (STS) marker, which was used to prepare a physical map of the Zmax1 (LRP5) gene
                                                                                      Sequence tagged site #235 used to prepare Zmax1 (LRP5) gene region map.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            New nucleic acid comprising a mutation in LRP5 or LRP6, useful for diagnosing a HBM-like phenotype in a subject and for preparing a composition for modulating bone mass and/or lipid levels in a subject suffering from e.g. osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                     Allen K, Anisowicz A, Graham JR, Morales A, Yaworsky PJ, Liu W;
                                                                                                                   Osteopathic, Gene therapy; High Bone Mass; HBM; LRPS; Zmaxl; LRP6; bone mass modulation; osteoporosis; STS; sequence tagged site; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; PRO polypeptide; secreted protein; transmembrane protein; cell death; neuropathy; neuropathy related disease; Charcot Marie-Tooth disorder; Refewn's disease; Krabbe's disease; chromosome mapping; gene mapping; genetic disorder; septic shock;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
2.9%; Score 12.2; DB 1; Length 18;
Best Local Similarity 82.4%; Pred. No. 4.7e+02;
Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 BP; 3 A; 8 C; 3 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                            GENOME THERAPEUTICS CORP.
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   ADB98354 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 GGCCAGGAGTGAACTG 17
                                                                                                                                                                                                                                                                                  11-MAY-2001; 2001US-0290071P.
17-MAY-2001; 2001US-029111P.
01-FEB-2002; 2002US-0353058P.
04-MAR-2002; 2002US-0361293P.
                                                                                                                                                                                                                                                       13-MAY-2002; 2002WO-US014877.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18 GGCAGGAGTGACTCTG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADB74025 standard; DNA; 18
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                                                           (first entry)
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                                                                                                                                                                                              WO200292000-A2
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                                                                                                                                                                  Homo sapiens,
                                                           04-DEC-2003
                                                                                                                                                                                                                         21-NOV-2002
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                               ADB98354;
                                                                                                                                                                                                                                                                                                                                                            (GENO-)
                                                                                                                                                                                                                                                                                                                                                                          (AMHP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 876
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XX
XX AC ADB7
XX
XX ADB7
XX ADB7
XX Huma
XX Huma
XX Huma
XX Huma
XX KW Cell
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ss; hepatitis C virus; HCV; HCV-associated disease; HCV infection;
fulminant hepatitis; chronic active hepatitis; cirrhosis;
hepatocellular carcinoma; cancer; tumour; lower relapse rate; antisense.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.9%; Score 12.2; DB 1; Length 18; 82.4%; Pred. No. 4.7e+02;
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             30-DEC-1999) 99W0-US031274-
06-JAN-2000) 2000W0-US000376-
11-FEB-2000) 2000W0-US000376-
11-FEB-2000) 2000W0-US000376-
11-FEB-2000) 2000W0-US000376-
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11-FEB-2000) 2000W0-US000376-
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11-MAR-2000) 2000W0-US016439-
11-MAR-2000) 2000W0-US016439-
11-MAR-2000) 2000W0-US016439-
11-MAR-2000) 2000W0-US014941-
22-MAR-2000) 2000W0-US014941-
02-JUL-2000) 2000W0-US014941-
02-JUL-2000) 2000W0-US014941-
02-JUL-2000) 2000W0-US014941-
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01-DEC-2000) 2000W0-US014965-
22-MAR-2001) 2001W0-US014956-
22-MAR-2001) 2001W0-US017090-
10-MAY-2001) 2001W0-US01700-
10-MAY-2001) 2001W0-US01700-
10-MAY-2001) 2001W0-US01700-
10-MAY-2001) 2001W0-US01700-
10-MAY-20010-
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10-MAY-20010-
10-MAY-20010-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 82.4
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GETH ) GENENTECH INC
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98US-0084414P.
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98US-0084627P.
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98US-0084637P.
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98US-008582P.
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98US-0085813P.
98US-00858173P.
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98US-013268P.
98US-0131848P.
99US-0131848P.
99US-0131822P.
99US-0131848P.
99US-0131848P.
99US-0131848P.
99US-0131848P.
99US-0131848P.
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15-MAY-1998 15-MAY-1998 16-MAY-1998 18-MAY-1998 22-MAY-1998 22-MAY-1998 22-MAY-1998 22-MAY-1998 28-MAY-1998 28-MAY-1998 26-JUN-1998 26-JUN-1998 30-JUL-1998 11-SEP-1998 11-SEP-1998 07-OCT-1998 07-OCT-1998 06-NOV-1998 26-NOV-1998

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Gaps

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Indels

23-UUN-1999 23-UUN-1999 26-UUL-1999 26-UUL-1999 25-AUG-1999 25-AUG-1999 25-AUG-1999 25-AUG-1999 25-AUG-1999 25-AUG-1999

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The invention relates to a new hepatitis C virus (HCV) genomic or messenger RNA antisense oligonuclectide. The oligonuclectide is useful disease. For preventing an HCV-associated disease, e.g., HCV infection, fulminant hepatitis, chronic active hepatitis, cirrhosis or hepatocellular carcinoma. Also for detection of HCV, HCV infection and HCV associated diseases. The oligonuclectide gives a more effective treatment than interferon alone with lower relapse rates. The present sequence represents a HCV antisense oligonuclectide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ss; primer; virucide; anti-HIV; hepatotropic; antiinflammatory; cytostatic; vulnerary; antiasthmatic; antiallergic; dermatological; antidiabetic; immunosuppressive; antirheumatic; antiarthritic; thyromimetic; protozoacide; amoebicide; antibacterial; gene therapy; virus; viral infections; non-viral infections; proliferative disease; inflammatory disease; allergic disease; autoimmune disease; PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vaccinia lister/PPVO NZ2 recombinant VVOV 96 PCR primer PPVO22r-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 New oligonucleotide, useful for preparing a composition for detect
treating or preventing HCV-associated disease, e.g. HCV infection,
fulminant hepatitis, chronic active hepatitis, cirrhosis or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.9%; Score 12.2; DB 1; Length 18; 82.4%; Pred. No. 4.7e+02; ative 0; Mismatches 3; Indels
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ADB79744/c
ID ADB79744 standard; DNA; 18 BP.
                                                                                                                                     93WO-JP001293.
95US-00397220.
95US-00452841.
96US-00650093.
97US-0098321.
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                                                                                      2001US-00853409
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hepatocellular carcinoma.
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Matches 14; Conservative
                                                                                                                                                                                                                                                                  (ANDE/) ANDERSON K P.
(HANE/) HANEGAK R C.
(NOZA/) NOZAKI C.
(DORR/) DORR F A.
(KWOH/) KWOH T J.
                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-697202/66
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               US2003171313-A1
                                                                                                                     10-SEP-1992;
10-SEP-1993;
09-MAR-1995;
30-MAY-1995;
17-MAY-1996;
10-DEC-1997;
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                                                                                    11-MAY-2001;
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                                                    11-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a new hepatitis C virus (HCV) genomic or messenger RNA antisense oligonucleotide. The oligonucleotide is useful disease. For preparing a composition for treating or preventing an HCV-associated disease, e.g., HCV infection, fulminant hepatitis, chronic active hepatitis, cirrhosis or hepatocellular carcinoma. Also for detection of HCV HCV infection and HCV associated diseases. The oligonucleotide gives a more effective treatment than interferon alone with lower relapse rates. The present sequence represents a HCV antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ss; hepatitis C virus; HCV; HCV-associated disease; HCV infection; fulminant hepatitis; chronic active hepatitis; cirrhosis; hepatocellular carcinoma; cancer; tumour; lower relapse rate; antisense.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           New oligonucleotide, useful for preparing a composition for detecting, treating or preventing HCV-associated disease, e.g. HCV infection, fullminant hepatitis, chronic active hepatitis, cirrhosis or hepatocellular carcinoma.
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2.9%; Score 12.2; DB 1; Length 18;
Best Local Similarity 82.4%; Pred. No. 4.7e+02;
Matches 14; Conservative 0; Mismatches 3; Indels
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/*tag= a
/mod_base= OTHER
/note= "OTHER = phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                      Kwoh TJ;
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                                                                                                                       10-SEP-1992; 92US-00945289.

10-SEP-1993; 93W0-7001293.

09-MAR-1995; 95US-00397220.

30-MAY-1996; 95US-00452841.

17-MAY-1996; 96US-00650093.

10-DEC-1997; 97US-0098321.

18-OCT-2000; 2000US-00690936.
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                                                                                          11-MAY-2001; 2001US-00853409
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                                                                                                                                                                                                                                                                          ANDERSON K P.
                                                                                                                                                                                                                                                                                            HANECAK R C.
NOZAKI C.
DORR F A.
KWOH T J.
                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-697202/66
                   US2003171313-A1.
                                                      11-SEP-2003
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Gaps

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Kwoh TJ;

Dorr FA,

(HANE/) (NOZA/) (DORR/) (ANDE/)

KWOH/)

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Wed Apr 21 12:58:21 2004
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New polynuclectide and recombinant proteins of Parapoxvirus ovis, useful for manufacturing a medicament for treating virus related disease, viral infections, non-viral infections, proliferative disease or inflammatory
                                                      Friederichs SM, Siegling A, Schlapp T, Mercer AA;
             L2-JUN-2002; 2002WO-EP006440.
                           13-JUN-2001; 2001NZ-00512341.
                                                                          WPI; 2003-221750/21
                                        (FARB ) BAYER AG
                                                 Weber O, Fr.
23-JAN-2003
                                                                                                               disease.
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Query Match
Best Local Similarity 82.4%;
Matches 14; Conservative

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ADB54025; #X#X#X8X#X#X#X#X#X#X#

Maier S, 27-FEB-2002; 2002EP-00004551. EPIG-) EPIGENOMICS AG P, Burger M, Schmitt A; Adorjan P, Rujan T,

Detecting and differentiating between colon cell proliferative disorders associated with a gene or its regulatory regions comprises contacting a target nucleic acid in a biological sample obtained from the subject with WPI; 2003-731620/69.

Nimmrich I, Becker E, Lesche R;

Claim 39; SEQ ID NO 81; 74pp; English.

a reagent

The invention relates to a novel method for detecting and differentiating between colon cell proliferative disorders associated with at least one cheween colon cell proliferative disorders associated with at least one reagent or regions. The method comprises contacting a target contacting of the contacting and target nucleic acid. The molecules of the invention demonstrate cytostatic activity while the method may useful contacts including cancers such as colon cell proliferative disorders, including cancers such as colon acenoma and colon carcinoma. The PNA (peptide nucleic acid)-oligomers are useful as probes for determining cytosine methylation state or single nucleotide of the polymorphisms. The current sequence is that of the oligomometic of the cinvention which was used to analyse the CGG positions within the genomic CMA taken from Wipoweb.

Sequence 18 BP; 4 A; 7 C; 5 G; 2 T; 0 U; 0 Other;

°, Query Match
2.9%; Score 12.2; DB 1; Length 18;
Best Local Similarity 82.4%; Pred. No. 4.7e+02;
Matches 14; Conservative 0; Mismatches 3; Indels

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177 GAGTCCAAGGCACATAT 193 dagreceddeacar 18 ద ઠે

741/c ADB76741 standard; DNA; 18 ADB76741; RESULT 881 ADB76741/c

Human PRO DNA PCR primer #218. (first entry) 04-DEC-2003

Human; PRO polypeptide; secreted protein; transmembrane protein; cell death; neuropathy; neuropathy related disease; charact-Marie-Tooth disorder; Refsum's disease; Krabbe's disease; chromosome mapping; gene mapping; gene matibacterial; immunosuppressive; neuroprotective; PCR; primer; se

Homo sapiens.

US2003083248-A1.

16-OCT-2001; 2001US-00978757 01-MAY-2003

97US-0062250P. 97US-0064249P. 97US-0065311P. 97US-0066364P. 17-OCT-1997; 03-NOV-1997; 13-NOV-1997; 21-NOV-1997;

Example 1; Page 23; 51pp; English.

The invention relates to a novel purified and isolated polynucleotide (NI) of Parapoxvirus ovis (PPVO) comprising a nucleotide sequence (SI, not defined in the specification), or its complementary sequence, fragment or functional variant. A polynucleotide of the invention has virucide, anti-MIV, hepatotropic, antiinflammatory, cytostatic, virucide, antiasthmatic, antiallergic, dermatorlogical, antidabetic, immunosuppressive, antitheumatic, antiathritic, thyrominetic, protozoacide, ancebicide, and antibacterial activity. The polynucleotides may have a use in gene therapy. The recombinant proteins encoded by the polynucleotides, or recombinant viruses comprising a Vaccinia virus genome and fragments of a pPVO genome are useful for manufacturing plarmaceutical compositions for treating virus related disease (e.g. hepatitis, papillomatosis, herpes virus infections, liver fibrosis, HIV infections with mycobacteria, mycoplasma, anceba or plasmodia), proliferative disease (e.g. cancer, leukaemia, warts or other skin neoplasms), inflammatory disease (e.g. crohn's disease, COPD, asthma or conditions related to healing of wounds), allergic disease, copp, asthma or conditions related to healing of wounds), allergic disease, copp, asthma or conditions related to healing of wounds), allergic disease, the and/or conditions thyroidatis, rheumatofa arthritis or diabetes mellitus). The present sequence is used in the exemplification of the invention.

.. 0 Score 12.2; DB 1; Length 18; Pred. No. 4.7e+02; 0; Mismatches 3; Indels

37 ACGAAGATGGCCACCAC 53

ADB54025 standard; DNA; 18 BP

(first entry)

04-DEC-2003

Oligonucleotide 17 used to analyse CpG positions within genomic DNA.

colon cell proliferative disorder, non methylated CpG dinucleotide, cytostatic, cancer, adenoma, carcinoma, cytosine methylation state, 88.

WO2003072821-A2.

04-SEP-2003

27-FEB-2003; 2003WO-EP002035

Sequence 18 BP; 2 A; 4 C; 8 G; 4 T; 0 U; 0 Other;

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Gaps

17 ACGTACATGGCCACCGC 1

12-006-209-209-209-209-209-209-209-209-209-209	. თ თ თ თ	1000	n on	on .	on o	1 01	O1	0, 0	J. O	,, 0	, 0,																••						, 20	500	200	200	20		202	200		56	200		200	Ñ	Ñ		Ñ	7	200		-	
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11- MAR-1998; 12- MAR-1998; 12- MAR-1998; 20- MAR-1998; 20- MAR-1998; 20- MAR-1998; 20- MAR-1998; 20- MAR-1998; 21- MAR-1998; 22- MAR-1998; 22- MAR-1998; 31- MAR-1998; 32- MAR-1998; 33- MAR-1998; 33- MAR-1998; 34- MAR-1998; 35- MAR-1998; 36- MAR-1998; 37- MAR-1998; 38- MAR-1998; 38- MAR-1998; 39- MAR-1998; 39- MAR-1998; 39- MAR-1998; 39- MAR-1998; 39- MAR-1998; 39- MAR-1998; 30- MAR-1998; 30- MAR-1998; 30- MAR-1998; 30- MAR-1998; 31- MAR-1998;																																																						
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The present invention relates to the isolation of novel human PRO polypeptides are secreted and transmembrane proteins. The PRO polypeptides are secreted and transmembrane proteins. The PRO polypeptides are useful for detecting other PRO polypeptides, for linking bioactive molecules to cells expressing PRO polypeptides, for modulating biological activities of cells expressing PRO polypeptides, and for identifying agonists or antagonists. The bioactive molecule maybe a toxin, radiolabel or antibody, and cause cell death. The PRO polypeptides are useful for treating neuropathy and neuropathy related diseases such as Charcot-Marie-Tooth disorder, Refsum's disease, and Krabbe's disease. The polymucleotide sequences encoding and gene mapping, in the generation of antisense RNA and DNA, in the preparation of PRO polypeptides, for
                   Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL; Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME; Goddard A, Godowski PV, Grimaldi UC, Gurney AL, Hillan KJ; Kljavin LJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL; Stewart TA, Tumas D, Williams PM, Wood WI;
                                                                                                                                                                                                                        New PRO polypeptides useful for treating peripheral neuropathy, neuropathies associated with systemic disease such as post-polio syndrome or AIDS-associated syndrome.
                                                                                                                                                                                                                                                                                                                                      Example 95; Page 174; 425pp; English.
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Gaps
Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels
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RESULT 882 ADC44167/c ID ADC44167 standard; DNA; 18 ADC44167;

BP

18-DEC-2003 (first entry)

Human PRO 298 PCR primer #3

Human, ss. PCR; secreted protein, transmembrane protein, PRO, cytostatic, ophthalmological, antiarchritic, osteopathic, antirheumatic, vulnerary; auditory, tumour growth, retinal disorder; sports-related joint problem, articular cartilage defects, osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.

Homo sapiens.

US2003054986-A1

20-MAR-2003

16-OCT-2001; 2001US-00981915.

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30-JUL-1998

30-JUL-1999

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Human, 88; PCR, secreted protein, transmembrane protein, PRO, cytostatic, ophthalmological, antiarthritic, osteopathic, antirheumatic, vulnerary, auditory, tumour growth, retinal disorder, sports-related joint problem, articular cartilage defects, osteoarthritis, rheumatoid arthritis, wound healing, hearing loss, primer.
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Matches 14; Conservative 0; Mismatches 3; Indels
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(GETH) GENENTECH INC

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                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                  Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Baton DL;
                                                                                                                                                                                                                                                                                  Query Match
2.9%; Score 12.2; DB 1; Length 18;
Best Local Similarity 82.4%; Pred. No. 4.7e+02;
Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                   149 GGAGGCCGGCTTCGACT 165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97US-0062250P.
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01-UTN-2001; 2001US-00872035.
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14-UTN-2001; 2001US-00882636.
19-UTN-2001; 2001US-00882636.
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20-UTN-2001; 2001WO-US019692.
29-UTN-2001; 2001WO-US0217755.
30-UTL-2001; 2001US-00918885.
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                                                                                                                                                                                                                                                                                                                                                                                         GGAGGTCGACTTCCACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC63891/c
ID ADC63891 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                  (GETH ) GENENTECH INC
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30-MAR-1998;
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강 원

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PR 07-0CT-1994) 990K-1051441

PR 20-NOV-1994) 991K-01184116

PR 20-NOV-1994) 991K-01184116

PR 22-DEC-1994) 991K-0119344

PR 22-DEC-1994) 991K-0119314

PR 22-DEC-1994) 991K-0101817

PR 22-DEC-1994) 991K-0101817

PR 12-MR-1994) 991K-0101817

PR 12-MR-1994) 991K-01116218

PR 12-MR-1994) 991K-010166-1994

PR 12-MR-1994) 991K-01016279

PR 12-MR-1994) 991K-0101322

PR 2-MR-1994) 991K-0101322

PR 2-MR-2004) 2000K-01800134

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PR 2-MR-2004) 2000K-01800134

PR 2-MR-2004) 2000K-01800136

PR 2-MR-2004) 2000K-0180139

PR 2-MR-2004) 2000K-0180139
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                                                                                                                                                                                                                                                                             vulnerary; virucide; neuroprotective; cytostatic; gene therapy; tumour cell proliferation inhibitor; secereted and transmembrane protein; PRO; viral infection; wound healing; tissue growth; muscle eneration; muscle regeneration; amyotrophic lateral sclerosis; neuropathy; AlDS-associated neuropathy; diabetic peripheral neuropathy; chromosome identification; antagonist; tissue typing; immunohistochemical staining; primer; 88.
                                                                                                   Gaps
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                                                                                                   3; Indels
                                                                              Score 12.2; DB 1;
Pred. No. 4.7e+02;
0; Mismatches 3;
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98US-0079664P.
                                                                                                                         149 GGAGGCCGCTTCGACT 165
20-JUN-2001; 2001WO-US019692.
29-JUN-2001; 2001WO-US021066.
90-JUL-2001; 2001WO-US021735.
30-JUL-2001; 2001US-00918585.
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                                                                               Query Match
Best Local Similarity 82.4%;
Matches 14; Conservative
                                                                                                                                                                                                                                                            Human PRO 298 PCR primer #3.
                                                                                                                                    17 GGAGGTCGACTTCCACT
                                                                                                                                                                                                ADC66991 standard; DNA; 18
                                                                                                                                                                                                                                        18-DEC-2003 (first entry)
                                                    (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                      JS2003060406-A1.
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12. MAR. 1998;
13. MAR. 1998;
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07-OCT-1998;
07-OCT-1998;
02-NOV-1998;
06-NOV-1998;
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Example 95; SEQ ID NO 519; 472pp; English.

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18-FEB-2000; 2000MO-US0064341.
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2000WO-US000376.
2000WO-US003565.
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99WO-US031274.
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30-NOV-1999;
02-DEC-1999;
                                                                                  12-MAR-1999;
12-APR-1999;
14-MAY-1999;
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(GETH) GENENTECH INC

Ľ KP, Botstein D, Desnoyers L, Eaton DL, E. Fong S, Gao W, Gerber H, Gerritsen ME; PJ, Grimaldi UC, Gurney AL, Hillan KJ; Milliams PM, Wood WI; Roy MA, Shelton Williams PM, Wood WI; Ashkenazi AJ, Baker KP, B Ferrara N, Filvaroff E, F Goddard A, Godowski PJ, G Kljavin IJ, Rob SS, Napie Stewart TA, Tumas D, Will

WPI; 2003-596568/56.

Novel secreted and transmembrane polypeptides and polynucleotides encoding them, useful for treating wound healing, tissue growth and muscle generation and regeneration, amyotrophic lateral sclerosis or neuropathy.

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The invention describes an isolated secereted and transmembrane PRO
C polypeptide (I). PRO polypeptide such as PRO213, PRO700, PRO320 or PRO615
CC is useful in biotechnological and medical research, as well as in various
industrial applications. PRO polypeptide such as PRO300, PRO300,
PRO320, PRO320, PRO351, PRO352, PRO981, PRO615, PRO618, PRO703,
CC PRO860 or PRO864 is useful for therapeutic purposes. PRO503 is useful
therapeutically in vivo for lessening the effects of viral infection.
CC PRO800 useful for the treatment of wound healing, tissue growth and
therapeutically in vivo for lessening the effects of viral infection.
CC PRO300 is useful for the treatment of wound healing, tissue growth and
muscle generation and regeneration. PRO337 is useful for treating
CC muscle generation are repeated or word the sale of viral infection.
CC weeful for generating transgenic animals which are
cuseful for generating transgenic animals which are
useful for generating sequences, and so construct hybridisation probes for
capping the gene which encodes the PRO and for the genetic analysis of
individuals with genetic disorders, for recombinantly expressing (I) and
for chromosome identification. (I) is useful as molecular marker for
correcting rescreening compounds to identify those that mind: the PRO
cuseful for screening compounds to identify those that mind: (I) is also
cuseful for screening compounds to identify those that mind: (I) is also
cuseful for munohistochemical staining and/or assays for PRO en
fluids. Anti-PRO antibodies are useful in diagnostic assays for PRO en
efficity purification of PRO from recombinant cell culture or natural
caffinity purification of PRO from recombinant cell culture or natural
caffinity purification of PRO from recombinant cell out transmembrane PRO
cy
protein associated primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 4 A; 6 C; 5 G; 3 T; 0 U; 0 Other;
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2.9%; Score 12.2; DB 1; Length 18; 82.4%; Pred. No. 4.7e+02; ative 0; Mismatches 3; Indels 149 GGAGGCCGGCTTCGACT 165 ADC69115 standard; DNA; 18 BP. 17 GGAGGTCGACTTCCACT 1 Local Similarity 82.4 RESULT 886 Matches ADC69115, à 요

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Gaps ö

> Human PRO 298 PCR primer #3. 18-DEC-2003 (first entry) ADC69115;

Human, ss, PCR, secreted protein, transmembrane protein, PRO, cytostatic, ophthalmological, antiarthritic, osteopathic; antirheumatic; vulnerary; auditory, tumour growth, retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer. 97US-0062250P. 97US-0064249P. 97US-0065311P. 97US-0066364P. 98US-0077450P. 98US-0077632P. 24-OCT-2001; 2001US-00999834 JS2003064407-A1. Homo sapiens. 03-NOV-1997; 13-NOV-1997; 21-NOV-1997; 10-MAR-1998; 11-MAR-1998; 03-APR-2003. 17-OCT-1997;

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PR 111-MAR-1998 9805-0077641P
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     Human; ss; PCR; secreted protein; transmembrane protein; PRO; cytostatic; ophthalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth; retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels
24-AUG-2000, 2000WO-USO23328.
08-NOV-2000; 2000US-00709238.
27-NOV-2000; 2000US-0073249.
01-DEC-2000; 2000WG-USO32678.
20-DEC-2000; 2000WG-USO32678.
20-DEC-2000; 2000WG-USO32678.
22-MAR-2001; 2001WG-USO36520.
22-MAR-2001; 2001WG-USO366520.
22-MAR-2001; 2001WG-USO366520.
22-MAR-2001; 2001WG-USO366520.
10-MAY-2001; 2001WG-USO36620.
10-MAY-2001; 2001WG-USO36620.
10-MAY-2001; 2001WG-USO36620.
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10-MAY-2001; 2001WG-USO36620.
11-MUN-2001; 2001WG-USO3636.
11-MUN-2001; 2001WG-USO3636.
11-MUN-2001; 2001WG-USO3636.
11-MUN-2001; 2001WG-USO36692.
20-JUN-2001; 2001WG-USO36692.
20-JUN-2001; 2001WG-USO36692.
20-JUN-2001; 2001WG-USO3666.
09-JUL-2001; 2001WG-USO31735.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97US-0062250P.
97US-0064249P.
97US-0065311P.
98US-0077450P.
98US-0077641P.
98US-0077641P.
98US-0077641P.
98US-0077641P.
98US-0078041P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 887

ADC63175/C

XX

AC ADC63175;
XX

I 8-DEC-2003 (first entry)
XX

Human PRO 298 PCR primer #3.
XX

Hombilancical; antiarthrit:
XM

Articular cartilage defects; dry
XX

Homo sapiens.
XX

US2003068648-A1.
XX

NX

10-APR-2003.
XX

YX

10-APR-2003.
XX

XX

XX

XX

XX

10-APR-2003.
XX

NX

11-APR-1997; 97US-00613921.
XX

XX

XX

XX

11-APR-1999; 98US-007641P.
PR 11-MAR-1998; 98US-0077641P.
PR 11-MAR-1998; 98US-0077641P.
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PR 11-MAR-1998; 98US-0077641P.
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PR 20-MAR-1998; 98US-0077641P.
PR 20-MAR-1998; 98US-0078910P.
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8 6

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1000; 2000WC-US014042.

1000; 2000WC-US014941.

1000; 2000WC-US01264.

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1001; 2001WC-US09520.

1001; 2001WC-US017092.

1001; 2001WC-US017092.
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99US-0130232P
99US-0131022P
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99WO-US012252
99WO-US012552
99WO-US012552
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2000WO-US013705.
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98US-0113296F.
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2000WO-US000219.
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2000WO-US003565.
2000WO-US004341.
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                                                                                                                                                                                  16-JUN-1999;
30-NOV-1999;
02-DEC-1999;
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30-DEC-1999;
06-JAN-2000;
06-JAN-2000;
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11-FEB-2000;
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24-FEB-2000;
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28-APR-1999;
14-MAY-1999;
14-MAY-1999;
02-JUN-1999;
             28-MAY-1998
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06-JAN-1999
10-MAR-1999
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29-MAR-1999;
21-APR-1999;
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GETH) GENENTECH INC.

Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL; Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME; Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ; Kljavin IJ, Kwo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL; Stewart TA, Tumas D, Williams PM, Wood WI;

WPI; 2003-695924/66.

New isolated secreted and transmembrane PRO polypeptides, useful in the preparation of a medicament for treating a condition responsive to the polypeptide, and as therapeutic agents e.g. vaccines.

Example 95; SEQ ID NO 519; 467pp; English.

```
The invention relates to an isolated PRO polypeptide (secreted or transmembrane protein) having at least 80% amino acid sequence identity co an amino acid sequence chosen from 94 fully defined sequences as given in the specification (including PRO lacking its associated signal peptide, a PRO extracellular domain with or without its associated signal comprising to peptide. Also included are mucleic acids encoding the PRO proteins comprising the vector comprising a PRO nucleic acid), a host cell comprising the vector and producing PRO, a chimaeric molecule comprising PRO antibody. PRO337 polypeptide is useful for detecting a PRO493 polypeptide is useful for detecting a PRO493 polypeptide. Comprising PRO493 polypeptide is useful for detecting PRO315. PRO700 or PRO735 polypeptide is useful for detecting Compressive PRO725. PRO700 or PRO735 polypeptide is useful for detecting compressive molecule to a cell expressing PRO337 polypeptide. The bloacting Complexity PRO379 polypeptide is useful for linking a prodective molecule to a cell expressing PRO337 polypeptide. The bloactive molecule comprise causes death of the cell. PRO337 polypeptide is useful for linking a prodective molecule is the toxin, radiolabel, or an antibody. The bloactive molecule
                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; 88; PCR; secreted protein; transmembrane protein; PRO; cytostatic; ophthalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth, retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.
                                                                                                                                                                                                                                                                                                                                                                ö
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98US-0077643P.
98US-0077643P.
98US-00778936P.
98US-0078936P.
98US-0078936P.
98US-0078936P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC68240/c
ID ADC68240 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-OCT-2001; 2001US-00978423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human PRO 298 PCR primer #3
                                                                                                                                                                                                                                                                                                                                                                                                  149 GGAGGCCGGCTTCGACT
                                                                                                                                                                                                                                                                                                                                                                                                                  17 GGAGGTCGACTTCCACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-DEC-2003 (first entry)
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11-MAR-1998;
11-MAR-1998;
12-MAR-1998;
12-MAR-1998;
20-MAR-1998;
20-MAR-1998;
20-MAR-1998;
20-MAR-1998;
20-MAR-1998;
25-MAR-1998;
25-MAR-1998;
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27-MAR-1998;
27-MAR-1998;
27-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC68240;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 888
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rng.res

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12:58:21 2004
21
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98US-0079786P.
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2001WG-US01263. 07-OCT-1998;
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20-NOV-1998;
22-DBC-1998;
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12-MAR-1999;
12-MAR-1999;
12-MAR-1999;
14-MAY-1999;
14-MAY-1999;
16-UJN-1999;
17-MAR-2000;
18-FEB-2000;
19-UJN-2000;
19-UJN-2001;
19-UJ G_{CX}

(GETH) GENENTECH INC.

Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL; Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME; Goddard A, Goddwski PJ, Grimaldi JC, Gurney AL, Hillan KJ; Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton Stewart TA, Tumas D, Williams PM, Wood WI;

Ľ,

WPI; 2003-657582/62.

Novel secreted and transmembrane polypeptides, designated PRO polypeptides, and polynucleotides encoding them useful for treating kidney diseases, bone, cartilage and retinal disorders.

Example 95; SEQ ID NO 519; 468pp; English.

The invention relates to an isolated PRO polypeptide (secreted or transmembrane protein) having at least 80% amino acid sequence identity

```
to an amino acid sequence chosen from 94 fully defined sequences as given peptide. A Ro extracellular domain with or without its associated signal peptide). Also included are nucleic acids encoding the PRO proteins mentioned above, a vector comprising a PRO nucleic acid), a host cell comprising the vector and producing PRO, a chimaeric molecule comprising PRO fused to a heterologous amino acid sequence, and an anti-PRO antibody. PRO337 polypeptide is useful for detecting a PRO4993 polypeptide is a sample suspected of containing PRO4993 polypeptide. Similarly, PRO4993 polypeptide is useful for detecting PRO337 polypeptide, and PRO725, PRO700 or PRO739 polypeptide is useful for detecting PRO359 polypeptide, and PRO1559 polypeptide is useful for detecting PRO359 polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human, se; PCR; secreted protein, transmembrane protein; PRO; cytostatic; ophthalmological; antiarchritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth; retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.
                                                                                                                                                                                                             0; Gaps
                                                                                                                                                                                 Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels
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98US-0079668P
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98US-0079668P
98US-0079668P
98US-0079668P
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                                                                                                                                                                                                                                                                                                                             ADC41560/c
ID ADC41560 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                18-DEC-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US2003072745-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-OCT-1997;
03-NOV-1997;
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20-MAR-1998;
20-MAR-1998;
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27-MAR-1998;
27-MAR-1998;
27-MAR-1998;
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37-MAR-1998;
37-MAR-1998;
37-MAR-1998;
37-MAR-1998;
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31-MAR-1998;
31-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                       ADC41560;
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    88888888888888
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PR 01-APR-1998 | 98US-0080327P | PR 01-APR-1998 | 98US-0080337P | PR 01-APR-1998 | 98US-0080337P | PR 01-APR-1998 | 98US-0080334P | PR 08-APR-1998 | 98US-0081049P | PR 08-APR-1998 | 98US-0081071P | PR 08-APR-1998 | 98US-0081195P | PR 08-APR-1998 | 98US-0081195P | PR 08-APR-1998 | 98US-0081195P | PR 15-APR-1998 | 98US-0081195P | PR 15-APR-1998 | 98US-0081195P | PR 15-APR-1998 | 98US-0081195P | PR 22-APR-1998 | 98US-0081195P | PR 22-APR-1998 | 98US-0081195P | PR 22-APR-1998 | 98US-0081362P | PR 22-APR-1998 | 98US-0081392P |
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4 49 c

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PR 06-AWR-1999) 99WC-UR0050106.
PR 12-WR-1999) 99WC-UR0050100.
PR 12-WR-1999) 99WC-01203572P.
PR 21-AWR-1999) 99WC-01203572P.
PR 21-AWR-1999) 99WC-01203572P.
PR 21-AWR-1999) 99WC-01203273P.
PR 21-AWR-1999) 99WC-01302289.
PR 21-AWR-1999) 99WC-01302299.
PR 21-AWR-1999) 99WC-013029
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comprising the vector and producing PRO, a chimaeric molecule comprising PRO fused to a heterologous amino acid sequence, and an anti-PRO antibody. PRO337 polypeptide is useful for detecting a PRO4993 polypeptide in a sample suspected of containing PRO4993 polypeptide. Similarly, PRO4993 polypeptide is useful for detecting PRO337
                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                       .
0
                                                                                                                                               Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                    149 GGAGGCCGGCTTCGACT 165
|||||||||||||||||||||||17 GGAGGTCGACTTCCACT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              97US-0064249P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADC67615;
                                                                                                                                                                                                                                                                                                                                                                                  RESULT 890
ADC67615/c
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Wed Apr 21 12:58:21 2004
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rng.res

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99US-0126773P
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07-MAY-1998

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06-MAY-1998;
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99US-0130232P.
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99US-0131022P.
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99US-0145680P.
99US-0146680P.
99US-0145680P.
99US-0145680P. (GETH) GENENTECH INC WPI; 2003-743810/70. 26-APR 1999; 14-APR 1999; 14-APR 1999; 14-MAY 1999; 16-UN 1999; 23-UN-1999; 26-UUL-1999; 26-UUL-1999; 26-UUL-1999; 26-UUL-1999; 36-UUL-1999; 36-UUL-

Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerriteen ME;
Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napler MA, Pan J, Paoni NF, Roy MA, Shelton
Stewart TA, Tumas D, Williams PM, Wood WI;

Novel isolated secreted and transmembrane PRO polypeptides, useful in the preparation of a medicament for treating a condition responsive to the polypeptide, and as therapeutic agents e.g. vaccines.

Example 95; SEQ ID NO 519; 464pp; English.

The invention describes an isolated secereted and transmembrane PRO polypeptide (I). PRO polypeptide such as PRO213, PRO700, PRO320 or PRO615 is useful in biotechnological and medical research, as well as in various industrial applications. PRO polypeptide such as PRO300, PRO866, PRO703, PRO860, PRO310, PRO811, PRO352, PRO381, PRO615, PRO615, PRO615, PRO813, Is useful therapeutically in vivo for lessening the effects of viral infection. PRO200 is useful for the treament of wound healing, tissue growth and muscle generation and regeneration. PRO313 is useful for treating amyotrophic lateral sclerosis, neuropathy, AIDS-associated neuropathy or

Query Match

Length 18; Score 12.2; 2.98;

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wed Apr zi iz:58:zi z00%
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98US-0081955P

21-APR-1998; 21-APR-1998; 22-APR-1998; 22-APR-1998; 22-APR-1998; 23-APR-1998; 23-APR-1998; 29-APR-1998; 29-APR-1998;

29-APR-1998; 29-APR-1998; 29-APR-1998;

29-APR-1998

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                                                                 Gaps
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                                                                 Indels
. 4.7e+02;
                                                                 Mismatches
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                                  Pred.
0; Mis
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9705-0064349P.
9705-0065364P.
9705-0077641P.
9805-0077641P.
9805-0077641P.
9805-0077641P.
9805-0070891P.
9805-0078910P.
9805-0078910P.
9805-0078910P.
9805-0079658P.
9805-0079658P.
9805-0079639P.
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9805-0079639P.
9805-0080333P.
9805-0080333P.
9805-0081049P.
                                                                                                                              165
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-OCT-2001; 2001US-00978193
                              Best Local Similarity 82.4%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human PRO 298 PCR primer #3
                                                                                                                                  149 GGAGGCCGGCTTCGACT
                                                                                                                                                                                     17 GGAGGTCGACTTCCACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-DEC-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US2003073624-A1.
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11 - MAR - 1998

12 - WAR - 1998

13 - WAR - 1998

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31 - M
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08-APR-1998;
08-APR-1998;
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01-APR-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                        ADC62551;
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ADC62551/C

ADC62551/C

ACC ADC625

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ADC625

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ADC625

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ADC625

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ADC625

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98US-0082568P.
98US-0082704P.
98US-0082704P.
98US-0082704P.
98US-0082322P.
98US-0083322P.
98US-0083495P.
98US-0083495P.
98US-0083495P.
98US-0083495P.
98US-0083495P.
98US-0083564P.
98US-008359P.
98US-008359P.
98US-008359P.
98US-008359P.
98US-008359P.
98US-008359P.
98US-008359P.
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98US-008464P.
98US-0084649P.
98US-008569P.
98US-0086486P.
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98US-0086486P.
98US-0091316P.
98US-00184216.

07-MAY-1998; 07-MAY-1998;

13-MAY-1998; 13-MAY-1998; 15-MAY-1998; 15-MAY-1998; 15-MAY-1998; 15-MAY-1998;

15-MAY-1998; 15-MAY-1998; 15-MAY-1998;

15-MAY-1998; 18-MAY-1998; 22-MAY-1998;

29-APR-1998; 29-APR-1998; 30-APR-1998; 30-APR-1998; 06-MAY-1998; 07-MAY-1998; 07-MAY-1998; 07-MAY-1998; 07-MAY-1998; 07-MAY-1998; 07-MAY-1998; 98US-00202054 98US-0113296P-98US-0113296P-99US-0113621P-99US-01054465-99US-00254465-99US-00265686-99US-00265213-99US-01257313-99US-01257313-

> 05-JAN-1999; 05-MAR-1999; 08-MAR-1999;

01.7UL-1998 30.7UL-1998 11.5EP-1998 07.0CT-1998 07.0CT-1998 02.NOV-1998 06.NOV-1998 06.NOV-1998 07.DEC-1998 22.DEC-1998 22.DEC-1998 22.DEC-1998

22-MAY-1998 22-MAY-1998 22-MAY-1998 28-MAY-1998 28-MAY-1998 26-JUN-1998 26-JUN-1998 26-JUN-1998

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Human, ss, PCR, secreted protein, transmembrane protein, PRO, cytostatic, ophthalmological, antiarthritic, osteopathic, antirheumatic, vulnerary, auditory, tumour growth, retinal disorder, sports-related joint problem, articular cartilage defects, osteoarthritis, rheumatoid arthritis, wound healing, hearing loss, primer.
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97105 - 0064349P

97105 - 0065364P

98105 - 0077641P

98105 - 0077641P

98105 - 0077641P

98105 - 0077641P

98105 - 0077649P

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98105 - 0077691P

98105 - 0079658P

98105 - 0079658P

98105 - 0079658P

98105 - 0079658P

98105 - 0079663P

98105 - 0079728P

98105 - 0079738P

98105 - 00801334P

98105 - 00801334P

98105 - 00801334P

98105 - 00801334P

98105 - 00810170P

98105 - 00810170P
      RESULT 892
ADC42184 standard; DNA; 18 BP
XX
XX
ADC42184;
XX
XX
ADC42184;
XX
XX
XX
ADC42184;
XX
XX
Human PRO 298 PCR primer #3.
XX
W bythtalmological; antiarthrit.
XM auditory; tumour growth; retain XX
X US2003104998-A1.
XX
US2003104998-A1.
XX

C5-UNN-2003.
XX

C5-UNN-2003.
XX

C5-UNN-1997; 97US-0065250P.
RR
X1-NAR-1998; 98US-007450P.
RR
X1-NAR-1998; 98US-0077612P.
RR
X1-MAR-1998; 98US-007761P.
RR
X1-MAR-1998; 98US-007761P.
RR
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RR
X1-MAR-1998; 98US-007761P.
RR
X1-MAR-1998; 98US-0079329P.
RR
X1-MAR-1998; 98US-0079329P.
RR
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RR
X1-MAR-1998; 98US-0079329P.
RR
X1-MAR-1998; 98US-007932P.
RR
X1-MAR-1998; 98US-007932P.
RR
X1-MAR-1998; 98US-007932P.
RR
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RR
X1-MAR-1998; 98US-008107P.
RR
X1-MAR-1998; 98US-0081019P.
RR
X1-MAR-1998; 98US-008101P.
RR
X1-MAR-19
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               99US-0130232P

99US-0131032P

99US-01311832

99US-0131287P

99US-0134287P

99US-014568PP

99US-014968PP

99US-014968PP

99US-014968PP

99US-014968PP

99US-014968PP

99US-014969PP

99US-01496PP

99US
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Best Local Similarity 82.4%;
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-JAN-2000; 2
06-JAN-2000; 2
118-FEB-2000; 2
118-FEB-2000; 2
24-FEB-2000; 2
10-MAR-2000; 2
11-MAR-2000; 2
11-DEC-2000; 2
11-DEC-2000; 2
11-DEC-2000; 2
11-DEC-2000; 2
11-DEC-2000; 2
11-MAR-2001; 2
                                                                                                                                                                                                           07-JUL-1999;
26-JUL-1999;
26-JUL-1999;
25-AUG-1999;
25-AUG-1999;
25-AUG-1999;
30-DCT-1999;
02-DEC-1999;
16-DEC-1999;
16-DEC-1999;
30-DEC-1999;
30-DEC-1999;
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99US-00284291

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Gaps ; 0

GGAGGCCGGCTTCGACT 165

149 17

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GAGGICGACTICCACT

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98US-0082804P.
98US-0083736P.
98US-00833322P.
98US-00833322P.
98US-00833322P.
98US-0083496P.
98US-0083496P.
98US-008359P.
98US-0083554P.
98US-0083554P.
98US-0083554P.
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98US-0083554P.
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98US-0084568P.
98US-0084568P.
98US-0084568P.
98US-0085339P.
98US-008533P.
99US-01335P.
99US-01335P.
99US-01335P.
99US-01335P.
99US-013382.
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99WO-US010733.
99WO-US010735.
99WS-0140528.
99US-0146228.
99US-0146228.
99US-0146228.
99US-0146228.
99US-0146228.
99US-0146228.
99US-01380142.
99US-01380142.
99US-01380142.
99WO-US0380137.
99WO-US038058.
99WO-US038063.
99WO-US038063.
99WO-US038063.
99WO-US038063.
99WO-US038083.
99WO-US017980.
99WS-01US-00882436.
99WS-01US-00882436.
99WS-01US-00882436. (GETH) GENENTECH INC 14-MAY-1999;
16-UIN-1999;
23-UIN-1999;
26-UIL-1999;
28-UIL-1999;
28-UIL-1999;
28-UIL-1999;
29-OCT-1999;
30-UIN-1999;
30-UIL-1999;
30-UI

Query Match
2.9%; Score 12.2; DB 1;
Best Local Similarity 82.4%; Pred. No. 4.7e+02;
Matches 14; Conservative 0; Mismatches 3; 149 GGAGGCCGGCTTCGACT 165

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Gaps

; 0

Length 18; Indels

> 17

8 8

RESULT 893
ADD24791/c
ID ADD24791 standard; DN
XX
AC ADD24791;

BP

ss; primer; detection; beer-spoilage; lactic acid bacteria; Gram-negative bacteria; spoilage bacteria.

Lactobacillus coryniformis

WO2002103043-A2

27~DEC-2002

19-JUN-2001; 2001DE-01029410. 19-JUN-2002; 2002WO-EP006808

(VERM-) VERMICON AG.

Snaidr J;

Beimfohr C,

WPI; 2003-175243/17.

Beer spoilage-associated primer SEQ ID 256.

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This invention describes a novel diagnostic kit for determining tolerance of pharmaceuticals in humans by determining allelic variability of at least two polymcrphisms of a human enzyme involved in phase I and/or II of the decoxification mechanism in a blood, tissue or other human sample, where tolerance is determined from presence or absence of alleles. The kit comprises two pairs of oligonucleotide primers, in which each pair amplifies, by PCR, part of a gene for a human detoxification mechanism associated enzyme. The kit may also contain two further pairs of oligonucleotides, serving as probes for detection of amplified DNA segments, especially where the probes are complementary to a single strand of one allele of the target gene. The probes are labelled with fluorophores (LC-Red40 or LC-Red705 for 5'-labelling or fluorescein for 3'-labelling) which generate a different signal in the hybridized condition. The enzymes detected include NAT2, CYP2D6, CYP1A2, CYP3A4, mEH, TPMT, MTHFR, paraoxonase, CYP2C19, CYP2C19, CYP2D1 or DPD. The kit is used to determine an individual's tolerance of a particular drug, to establish a suitable dose and/or to predict if a subject will subject and reliable determination of the metabolic capacity of phase I and/or II enzymes at the molecular level. This sequence represents a probe used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ò
                                                                                  diagnostic; pharmaceutical tolerance; side effect; drug; human; allelic variability; polymorphism; phase I; phase II; drug; detoxification mechanism; PCR; primer; probe; NAT2; CYP2D6; CYP1A2; CYP3A4; mEH; TPWT; MTHRR; paracoxonase; CYP2C9; CYP2C19; CYP2E1; DPD; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Diagnostic kit, useful for assessing a subject's tolerance of drugs, comprises reagents for determining alleles of genes encoding detoxification enzymes.
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2.9%; Score 12.2; DB 1; Length 18;
Best Local Similarity 82.4%; Pred. No. 4.7e+02;
Matches 14; Conservative 0; Mismatches 3; Indels
                                           Human CYP2D6 mutants G1846A and G1758T probe H234.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18 BP; 2 A; 3 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Waschuetza S, Schnakenberg E, Lustig M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 6; Page 19; 156pp; German.
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                                                                                                                                                                                                                                                                                                                                                                 2001DE-01040651.
2002DE-01019373.
                                                                                                                                                                                                                                                                                                                          22-AUG-2002; 2002WO-EP009386.
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  15-JAN-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-290079/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                   (ADNA-) ADNAGEN AG
                                                                                                                                                                                                                                      WO2003018837-A2.
                                                                                                                                                                                                                                                                                                                                                                 24-AUG-2001;
30-APR-2002;
                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                       detecting beer-spoilage bacteria in sample. The bacteria detected include lactic acid bacteria in assumple. The bacteria detected include lactic acid bacteria of the genera Lactobacillus or Pediococcus, especially the species L. coryniformis, L. perolens, L. buchneri, L. plantarum, L. fructivorans, L. lindneri, L. casei, L. brevis or P. damnosus or Gram-negative bacteria of the genera Pectinatus and M. cerevisiae. The oligonucleocides of the invention provide rapid detection of spoilage bacteria (typically within 48 hours, compared with 7-12 days for conventional culture methods), can detect all relevant bacteria in parallel, can differentiate between species of the same genus, and are easy to use ADBISS247 represent the oligonucleotides used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                This invention describes novel oligonucleotides used in a method for
                                                                                                                                                                                                                                                                                                                      New oligonucleotides, useful for rapid detection of beer-spoilage
bacteria by in situ hybridization, are specific for type, genus or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88; primer; detection; beer-spoilage; lactic acid bacteria;
Gram-negative bacteria; spoilage bacteria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
2.9%; Score 12.2; DB 1; Length 18;
Best Local Similarity 82.4%; Pred. No. 4.7e+02;
Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18 BP; 3 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Beer spoilage-associated primer SEQ ID 262.
                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; SEQ ID NO 256; 88pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            103 CTGACCGCGACCGCAGC 119
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ADE15067 standard; DNA; 18 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nethod of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADE15067;
                                                                                                                                                                                                                                                                                                                                                        species.
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ADE15061 standard; DNA; 18 BP

(first entry)

29-JAN-2004

ADE15061; RESULT 894
ADE15061/C
ID ADE150
XX
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DT 29-JAN
XX

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9805-078910P-9805-078910P-9805-0778910P-9805-0778910P-9805-0778910P-9805-0778910P-9805-0778910P-9805-0778910P-9805-0779961P-9805-0779961P-9805-0779961P-9805-0779923P-9805-0779923P-9805-0779923P-9805-0779923P-9805-0779923P-9805-0779923P-9805-0779923P-9805-0779923P-9805-0779923P-9805-0779923P-9805-0779923P-9805-0779923P-9805-077992P-9805-077992P-9805-077992P-9805-077992P-9805-077992P-9805-077992P-9805-077992P-9805-077992P-9805-07799P-9805-077992P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-07799P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779P-9805-0779
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22-APR-1998;
27-APR-1998;
28-APR-1998;
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29-APR-1998;
66-MAY-1998;
66-MAY-1998;
66-MAY-1998;
67-MAY-1998;
67-MAY-1998;
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15-APR-1998;
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                  20-MAR-1998;
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31-MAR-1998;
31-MAR-1998;
31-MAR-1998;
31-MAR-1998;
31-MAR-1998;
31-MAR-1998;
31-MAR-1998;
31-APR-1998;
This invention describes novel oligonuclectides used in a method for detecting beer-spoilage bacteria in a sample. The bacteria detected include lactic acid bacteria of the genera Lactobacillus or Pediococcus, especially the species L. coryniformis, L. perolens, L. buchneri, L. plantarum, L. fructivorans, L. lindmeri, L. casel, L. brevis or P. damnosus or Gram-negative bacteria of the genera Pectinatus and Megasphaera, specifically P. frisingensis, P. cerevisiiphilus and M. cerevisiae. The oligonucleotides of the invention provide rapid detection of spoilage bacteria (typically within 48 hours, compared with 7-12 days for conventional culture methods), can detect all relevant bacteria in parallel, can differentiate between species of the same genus, and are method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human, ss; PCR; secreted protein; transmembrane protein; PRO; cytostatic; ophthalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth; retinal disorder; sports-related joint problem; articular cartilage defects; osteopathritis; rheumatoid arthritis; wound healing; hearing loss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
                                                                                                                                                                                              New oligonucleotides, useful for rapid detection of beer-spoilage
bacteria by in situ hybridization, are specific for type, genus or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match 2.9%; Score 12.2; DB 1; Length 18; Local Similarity 82.4%; Pred. No. 4.7e+02; es 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18 BP; 2 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID NO 262; 88pp; German.
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ADE49553/C
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AC
ADE49553;
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XY
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C 9-JAN-2004 (first entry)
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XX
Human PRO 298 PCR primer #3.
XX
XX
XX
Human, ss; PCR; secreted prote with auditory; tumour growth; retiring auditory; tumour growth; retiring articular cartilage defects; XX
XX
Homo sapiens.
XX
XX
CSC03096744-Al.
XX
XX
X2-MAY-2003.
XX
X2-MAY-2003.
XX
YX
Z2-MAY-2003.
XX
YX
Z2-MAY-2003.
XX
YX
X2-MAY-2003.
XX
XX
X2-MAY-2003.
XX
YX
X17-0CT-1997; 97US-006256P.
PR 11-NAR-1998; 98US-007764P.
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97US-0064249P.
97US-006331P.
97US-0077450P.
98US-0077632P.
98US-0077641P.
98US-0077641P.
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     19-JUN-2001; 2001DE-01029410.
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                                                  (VERM-) VERMICON AG
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Best Local Si
Matches 14
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2000US-00747259

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98US-0086392P.
98US-0086414P.
98US-0086418P.
98US-0087208P.
98US-0087208P.
98US-00105413.
98US-0010513P.
98US-0010518P.
98US-0010683P.
98US-0010683P.
98US-001683P.
98US-0018738P.
99US-001873P.
99US-00254465.
99US-00254465.
99US-001873P.
99US-001873P.
99US-001873P.
99US-013182P.
99US-013182P.
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99US-013182P.
99US-013183P.
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99US-013183P.
99US-0146222P.
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99US-0146222P.
99US-0146222P.
99US-0182885E.
99WO-US02885E.
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2000WO-US000277.
2000WO-US000376.
2000WO-US003565.
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06-JAN-2000;
11-FEB-2000;
24-FEB-2000;
02-MAR-2000;
21-MAR-2000;
31-MAR-2000;
32-MAY-2000;
32-MAY-2000;
32-MAY-2000;
32-MAY-2000;
32-MAY-2000;
32-MAY-2000;
32-MAY-2000;
32-MAY-2000;
32-MAY-2000;
33-MAY-2000;
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02-DEC-1999;
16-DEC-1999;
30-DEC-1999;
30-DEC-1999;
05-JAN-2000;
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02-JUN-1999
13-JUN-1999
07-JUL-1999
26-JUL-1999
25-JUL-1999
25-AUG-1999
25-AUG-1999
25-AUG-1999
25-AUG-1999
25-AUG-1999
30-NOV-1999
22-MAY-1998;

22-MAY-1998;

22-MAY-1998;

28-MAY-1998;

28-MAY-1998;

28-MAY-1998;

28-MAY-1998;

26-JUN-1998;

30-JUL-1998;

31-JUL-1999;

32-DEC-1999;

33-DEC-1999;

33-DEC-1999;

33-DEC-1999;

33-DEC-1999;

33-DEC-1999;

33-DEC-1999;

34-MAR-1999;

31-MAR-1999;

31
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Human; ss; PCR; secreted protein; transmembrane protein; PRO; cytostatic; ophthalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth, retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL; Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritesn MB; Goddard A, Goddowski PJ, Grimaldi JC, Gurney AL, Hillan KJ; Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton Stewart TA, Tumas D, Williams PM, Wood WI;
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Pred. No. 4.7e+02;
0; Mismatches 3;
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ADE35607/C

ID ADE35607 standard; DNA; 18 BP. XX

AC ADE35607;

XX

ALE35607;

XX

DT 29-JAN-2004 (first entry)

XX

Human PRO 298 PCR primer #3.

XX

Human, ss; PCR; secreted protein

XM

Auditory; tumour growth; retinal

XM

NS2003203434-A1.

XX

NS2003203434-A1.

XX

B-CCT-2001; 2001US-00145088.

XX

18-CCT-2001; 2001US-00145088.

XX

18-CR-1999; 99US-0131445P.

PR

18-MAY-1999; 99US-0131445P.

PR

25-AUG-1999; 99US-0131445P.

PR

25-AUG-1999; 99US-0131445P.

PR

25-AUG-1999; 99US-0131445P.

PR

25-AUG-1999; 99US-0131445P.

PR

30-UUL-2001; 2001US-0018585.

XX

Ashkenazi AJ, Baker KP, Botste

PI Ferrara N, Filvaroff E, Fong SP

PI Goddard A, Godowski PJ, Grimal

PI Kljavin IJ, Kuo SS, Napier MA,

XX

XX

WPI; 2003-875641/81.
          20-DEC-2000; 2000MO-US034956.

28-FRB-2001; 2001MO-US034956.

22-MAR-2001; 2001US-00816744.

22-MAR-2001; 2001US-00816744.

22-MAR-2001; 2001WC-008552.

10-MAY-2001; 2001WC-00854280.

10-MAY-2001; 2001WC-00854280.

25-MAY-2001; 2001WC-00817092.

01-JUN-2001; 2001WC-00817800.

05-JUN-2001; 2001WC-00817800.

19-JUN-2001; 2001WC-0081636.

19-JUN-2001; 2001WC-00886342.

20-JUN-2001; 2001WC-00886342.

20-JUN-2001; 2001WC-00886342.

20-JUN-2001; 2001WC-00886342.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   149 GGAGGCCGGCTTCGACT 165
17 GGAGGTCGACTTCCACT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 2.9%;
Best Local Similarity 82.4%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                     Ashkenazi AJ, Baker KP,
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New genes, and its encoded secreted and transmembrane polypeptides, useful for treatling e.g. lung or breast tumors, osteoarthritis, rheumatoid arthritis, obesity, diabetes, hyperinsulinemia, hypoinsulinemia or wounds.

Example 95; SEQ ID NO 519; 462pp; English.

The invention relates to an isolated PRO polypeptide (secreted or transmembrane protein) having at least 80% amino acid sequence identity con an amino acid sequence chosen from 94 fully defined sequences as given in the specification (including PRO lacking its associated signal peptide). Also extractabiliar domain with or without ties associated signal peptide). Also included are unclait a chids encoding the PRO proteins mentioned above, a vector comprising a pro unclaic acid) a host call septide). Also included are unclaid a chids encoding the PRO proteins mentioned above, a vector comprising a pro unclaic acid) a host call peptide. Subject of a heterologous amino acid sequence, and an anti-PRO comprising the vector and producing PRO a chidsactic molecule comprising the vector and producing PRO a chidsactic molecule comprising the vector and producing PRO a chidsactic molecule is the propertie is useful for detecting PRO plypeptide in a sample suspected of containing PRO4993 polypeptide. PRO4993 polypeptide is useful for detecting PRO4993 polypeptide is useful for detecting PRO4993 polypeptide. PRO4993 polypeptide is useful for linking a bioactive molecule to a cell expressing PRO4993 polypeptide; PRO4993 polypeptide is useful for linking a bioactive molecule is the toxin. radiolabel. Or an artibody. The bioactive molecule is the toxin. radiolabel. Or an artibody The bioactive molecule is useful for linking a bioactive molecule is useful for linking a bioactive molecule is useful for mining a bioactive or anti-PRO493 polypeptide is useful for mining the biological activity of the call expressing PRO4993 polypeptide is useful for modulating the biological activity of the call expressing PRO4993 polypeptide is useful for modulating the biological activity of the call expressing PRO4993 polypeptide is useful for minipate to manipage defects. The prosessing PRO4993 polypeptide is useful for minipage defects. The polypeptide is useful for minipage and profised activity of the cell expressing PRO4993 polypeptide is

Seguence 18 BP; 4 A; 6 C; 5 G; 3 T; 0 U; 0 Other;

2.9%; Score 12.2; DB 1; Length 18; 82.4%; Pred. No. 4.7e+02; vative 0; Mismatches 3; Indels 0; Gaps 14; Conservative Query Match Best Local Similarity Matches

ADE16721 standard; DNA; 18 BP ADE16721/c

ADE16721;

29-JAN-2004 (first entry)

Human PRO 298 PCR primer #3.

Human; ss; PCR; secreted protein; transmembrane protein; PRO; cytostatic; ophthalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth; retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.

Homo sapiens

US2003203435-A1.

30-OCT-2003

18-OCT-2001; 2001US-00145092

99WO-US005028. 99US-0141037P. 99US-00380138. 30-APR-1998; 08-MAR-1999;

23-JUN-1999; 99US-0141037P. 25-AUG-1999; 99US-00380138. 18-FEB-2000; 2000WO-US04341. 30-JUL-2001; 2001US-00918858.

(GETH) GENENTECH INC.

Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL; Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME; Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ; Kljavin IJ, KNO SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL; Stewart TA, Tumas D, Williams PM, Wood WI;

WPI; 2003-875642/81.

New genes, and its encoded secreted and transmembrane polypeptides, useful for treating e.g. lung or breast tumors, osteoarthritis, theumatoid arthritis, obsesty, diabetes, hyperinsulinemia, hypoinsulinemia or wounds.

Example 95; SEQ ID NO 519; 452pp; English.

The invention relates to an isolated PRO polypeptide (secreted or transmembrane protein) having at least 80% amino acid sequence identity to an amino acid sequence those from 94 fully defined sequences as given in the specification (including PRO lacking its associated signal opptide, a PRO extracellular domain with or without its associated signal peptide. Also included are nucleic acids encoding the PRO proteins mentioned above, a vector comprising a PRO nucleic acid), a host cell comprising the vector and producing PRO, a chimaeric molecule comprising the vector and producing PRO, a chimaeric molecule comprising prolypeptide is useful for detecting PRO337 polypeptide is useful for detecting PRO337 polypeptide is useful for detecting prolypeptide. PRO493 polypeptide is useful for distance. Detecting prolypeptide is useful for linking a bloactive molecule to a cell expressing PRO337 polypeptide; prolypeptide is useful for linking a bloactive molecule to a cell expressing PRO493 polypeptide; prolypeptide is useful for linking a bloactive molecule to a cell expressing PRO493 polypeptide; prolypeptide is useful for modulating at least one biological activity of the cell expressing PRO337 polypeptide is useful for modulating at least one biological activity of the cell expressing PRO393 polypeptide is useful for modulating the biological activity of the cell expressing PRO393 polypeptide is useful for modulating the biological activity of the cell expressing PRO393 polypeptide is useful for modulating the biological activity of the cell expressing PRO393 polypeptide is useful for modulating the biological activity of the cell expressing PRO393 polypeptide is useful for modulating the biological activity of the cell expressing PRO393 p acid encoding a PRO protein.

Sequence 18 BP; 4 A; 6 C; 5 G; 3 T; 0 U; 0 Other;

2.9%; Score 12.2; DB 1; Length 18; 82.4%; Pred. No. 4.7e+02; Query Match Best Local Similarity

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 Mismatches
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The invention relates to an isolated PRO polypeptide (secreted or transmembrane protein) having at least 80% amino acid sequence identity transmembrane protein) having at least 80% amino acid sequence chosen from 94 fully defined sequences as given in the specification (including PRO lacking its associated signal peptide, a PRO extracellular domain with or without its associated signal peptide. Also included are nucleic acids encoding the PRO proteins mentioned above, a vector comprising PRO, a chimaeric molecule comprising the vector and producing PRO, a chimaeric molecule comprising PRO fused to a heterologous amino acid sequence, and an anti-PRO antibody. PRO337 polypeptide is useful for detecting a PRO4993 polypeptide is a sample suspected of containing PRO4993 polypeptide.

Similarly, PRO4993 polypeptide is useful for detecting PRO37 polypeptide is useful for detecting PRO725, PRO700 or PRO759 polypeptide is useful for detecting PRO725, PRO700 or PRO4993 polypeptide is useful for letecting PRO725, PRO700 or PRO4993 polypeptide is useful for linking a bloactive molecule to a cell expressing PRO337 polypeptide. The bioactive Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL; Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME; Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ; Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL; Stewart TA, Tumas D, Williams PM, Wood WI; Human; ss; PCR; secreted protein; transmembrane protein, PRO; cytostatic; ophthalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth; retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer. New PRO genes and encoded secreted and transmembrane polypeptides, u for treating e.g. lung or breast tumors, osteoarthritis, rheumatoid arthritis, obesity, diabetes, hyperinsulinemia, hypoinsulinemia or Example 95; SEQ ID NO 519; 453pp; English. 165 98US-0086414P. 98US-0113296P. 98WO-US000106. 99WO-US005028. 99US-00284291. ADD73336 standard; DNA; 18 BP 18-OCT-2001; 2001US-00145129 18-FEB-2000; 2000WO-US004341 30-JUL-2001; 2001US-00918585 Н Human PRO 298 PCR primer #3. 149 GGAGGCCGGCTTCGACT 17 GGAGGTCGACTTCCACT 29-JAN-2004 (first entry) GETH) GENENTECH INC. WPI; 2003-875643/81. US2003203436-A1. Homo sapiens. 05-JAN-1999; 08-MAR-1999; 25-AUG-1999; 22-MAY-1998; 30-OCT-2003. 22-DEC-1998 12-APR-1999 wounds. RESULT 899 Matches ADD73336,

cc molecule is the toxin, radiolabel, or an antibody. The bioactive molecule by the cell. PR0337 polypeptide is useful for linking a bioactive molecule to a cell expressing PR04939 polypeptide; PR0755, PR0755, PR0750 or PR0739 polypeptide are useful for linking a bioactive molecule to a cell expressing PR0755, cto a cell expressing PR01559 polypeptide; and PR01559 polypeptide is useful for linking a bioactive molecule to a cell expressing PR0725, PR0700 or PR0739 polypeptide; PR04993 polypeptide or anti-PR0337 polypeptide is useful for modulating at least one biological activity of the cell expressing PR0349 polypeptide is useful for modulating the coll expressing PR0337 polypeptide or anti-PR04993 polypeptide; PR0725, cto PR0700 or PR0739 polypeptide or an anti-PR0559 polypeptide; PR0725, cto PR0700 or PR0739 polypeptide or an anti-PR0555, anti-PR0700 or anti-PR0755, anti-PR0755, anti-PR0759 polypeptide; profiged activity of the cell expressing PR0459 polypeptide is useful for modulating the biological activity of the cell expressing PR0755, pr0700 or PR0739 polypeptide is useful for modulating tumour growth, retinal disorders, polypeptides are useful for inhibiting tumour growth, retinal disorders, sports-related joint problems, articular cartilage defects, sports-related joint problems, articular cartilage defects, costcoarthritis or rheumatoid arthritis, wound hearing and hearing loss in mammals. The present sequence is a PCR primer used to isolate nucleic acid encoding a PRO protein. ö Human, ss; PCR; secreted protein; transmembrane protein; PRO; cytostatic; ophthalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth, retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer. Gaps ; 0 Query Match
2.9%; Score 12.2; DB 1; Length 18;
Best Local Similarity 82.4%; Pred. No. 4.7e+02;
Matches 14; Conservative 0; Mismatches 3; Indels Sequence 18 BP; 4 A; 6 C; 5 G; 3 T; 0 U; 0 Other; 149 GGAGGCCGGCTTCGACT 165 99WO-US005190. 99WO-US008313. 99WO-US010733. 99WO-US012252. 99US-00380138. 98US-0079920P. 98WO-US021141. 98WO-US024855. 99WO-US000106. 99WO-US005028. ADD72694 standard; DNA; 18 BP. 17 GGAGGTCGACTTCCACT 1 19-OCT-2001; 2001US-00164929 Human PRO 298 PCR primer #3. 29-JAN-2004 (first entry) US2003194781-A1. Homo sapiens. 05-JAN-1999; 08-MAR-1999; 10-MAR-1999; 15-APR-1999; 14-MAY-1999; 02-JUN-1999; 16-OCT-2003. 07-0CT-1998; 20-NOV-1998; 30-MAR-1998 ADD72694; ADD72694/c 요 \$ ò

99WO-US030095. 99WO-US031243. 99WO-US031274.

99WO-US028313 99WO-US028551

25-AUG-1999; 02-DEC-1999; 02-DEC-1999

05-JAN-2000; 206-JAN-2000; 206-JAN-2000; 206-JAN-2000; 207-2000; 2 20-JUN-2001; 22-MAR-2001;

2001WO-US019692 2001US-00918585 (GETH) GENENTECH INC. 09-JUL-2001;

2000WO-US013705. 2000WO-US014042. 2000WO-US014041. 2000WO-US020710. 2000WO-US02333. 2000WO-US0334956. 2001WO-US034956.

17.07.21

2001WO-US009552. 2001WO-US017092. 2001WO-US017800.

DĽ; Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
Goddowski PJ, Grimaldi JC, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton I
Stewart TA, Tumas D, Williams PM, Wood WI;

WPI; 2003-852598/79.

New secreted and transmembrane PRO nucleic acids and polypeptides, useful for stimulating the release of tumor necrosis factor alpha from human blood and stimulating the proliferation of differentiation of chondrocyte cells.

Example 95; SEQ ID NO 519; 462pp; English

The invention relates to an isolated PRO polypeptide (secreted or transmembrane protein) having at least 80% amino acid sequence identity cransmembrane protein) having at least 80% amino acid sequences as given in the specification (including PRO lacking its associated signal opeptide). Also included are nucleic acids enceding the PRO proteins comprising the vector comprising a PRO nucleic acid), a host cell comprising the vector and producing PRO, a chimaeric molecule comprising the vector and producing PRO, a chimaeric molecule comprising CC comprising the vector and producing a PRO nucleic acid), a host cell comprising the vector and producing a PRO nucleic acid), a host cell comprising the vector and producing a PRO nucleic acid), a host cell comprising the vector and producing a PRO antibody. PRO337 polypeptide is useful for detecting a PRO493 polypeptide in a sample anspected of containing PRO493 polypeptide. PRO725, PRO700 or PRO739 polypeptide is useful for detecting PRO4559 polypeptide, and PRO1559 polypeptide is useful for detecting PRO455, PRO700 or PRO739. PRO493 polypeptide is useful for linking a causes death of the cell expressing PRO493 polypeptide; The bioactive molecule to a cell expressing PRO493 polypeptide; PRO705, cuseful for linking a bioactive molecule to a cell expressing PRO493 polypeptide; PRO700 or PRO739 polypeptide are useful for linking a bioactive molecule to a cell expressing PRO493 polypeptide; PRO493 polypeptide is useful for linking a bioactive molecule for a cell expressing PRO4933 polypeptide; PRO4939 polypeptide or anti-PRO4939 polyp modulating the biological activity of the cell expressing PRO1559

ö Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL; Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME; Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ; Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL; Stewart TA, Tumas D, Williams PM, Wood WI; Human; ss; PCR; secreted protein; transmembrane protein; PRO; cytostatic; ophthalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth; retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer. Gaps ; 0 Query Match 2.9%; Score 12.2; DB 1; Length 18; Best Local Similarity 82.4%; Pred. No. 4.7e+02; Matches 14; Conservative 0; Mismatches 3; Indels Sequence 18 BP; 4 A; 6 C; 5 G; 3 T; 0 U; 0 Other 06-MAY-1998; 98US-0084414P. 22-DEC-1998; 98US-013296P. 08-JAN-1999; 99WG-US000106. 08-MAR-1999; 99US-00284291. 25-AUG-1999; 99US-00380138. 18-FEB-2000; 2000WG-US004341. 30-JUL-2001; 2001US-00918585. 165 H 18-OCT-2001; 2001US-00145016 acid encoding a PRO protein. Human PRO 298 PCR primer #3. 17 GGAGGTCGACTTCCACT 149 GGAGGCCGGCTTCGACT ADE17345 standard; DNA; 18 29-JAN-2004 (first entry) (GETH) GENENTECH INC. JS2003203433-A1. Homo sapiens. 30-OCT-2003. ADE17345; RESULT 8888888888 ઠે 셤

polypeptide; and PRO1559 polypeptide or anti-PRO725, anti-PRO700 or anti-PRO739 polypeptide is useful for modulating the biological activity of the cell expressing PRO725, PRO700 or PRO739 polypeptide. The oplypeptide are useful for inhibiting tumour growth, retinal disorders, sports-related joint problems, articular cartilage defects, osteoarthritis or rheumatoid arthritis, wound healing and hearing loss in mammals. The present sequence is a PCR primer used to isolate nucleic

Example 95; SEQ ID NO 519; 459pp; English.

New genes, and its encoded secreted and transmembrane polypeptides, useful for treating e.g. lung or breast tumors, osteoarthritis, rheumatoid arthritis, obesity, diabetes, hyperinsulinemia, hypoinsulinemia or wounds.

WPI; 2003-875640/81.

The invention relates to an isolated PRO polypeptide (secreted or transmembrane protein) having at least 80% amino acid sequence identity to an amino acid sequence chosen from 94 fully defined sequences as given

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in the specification (including PRO lacking its associated signal peptide, a PRO extracellular domain with or without its associated signal peptide, Also included are mucleic acids encoding the PRO proteins mentioned above, a vector comprising a PRO nucleic acid), a host cell comprising the vector and producing PRO, a chimaeria molecule comprising comprising the vector and producing PRO, a chimaeria molecule comprising antibody. PRO337 polypeptide is useful for detecting a PRO493 polypeptide is asmile suspected of containing PRO493 polypeptide. Similarly, PRO493 polypeptide is useful for detecting PRO1559 polypeptide, and PRO1559 polypeptide is useful for detecting PRO1559 polypeptide. PRO725, PRO700 or PRO159 polypeptide is useful for linking a bioactive molecule to a cell expressing PRO4937 polypeptide is useful for linking a bioactive molecule to a cell expressing PRO4937 polypeptide; PRO705, CC causes death of the cell. PRO337 polypeptide; PRO705, CC causes death of the cell. PRO337 polypeptide; PRO725, CC cuseful for linking a bioactive molecule to a cell expressing PRO4939 polypeptide; and PRO1559 polypeptide is useful for linking a bioactive molecule for a cell expressing PRO4939 polypeptide or anti-PRO493 polypeptide is useful for modulating the biological activity of the cell expressing PRO4939 polypeptide or anti-PRO4939 polypeptide or anti-PRO4939 polypeptide or anti-PRO4939 polypeptide; propoptide or anti-PRO4939 polypeptide or a
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Sequence 18 BP; 4 A; 6 C; 5 G; 3 T; 0 U; 0 Other;

0; Gaps Query Match
2.9%; Score 12.2; DB 1; Length 18;
Best Local Similarity 82.4%; Pred. No. 4.7e+02;
Matches 14; Conservative 0; Mismatches 3; Indels

149 GGAGGCCGGCTTCGACT 165

(first entry)

Homo sapiens.

US2003104536-A1.

05-JUN-2003

07-OCT-1998; 20-NOV-1998; 05-JAN-1999;

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17 GGAGGTCGACTTCCACT ઠે 셤

RESULT 902 ADE48853/c ID ADE48853 standard; DNA; 18 BP. 29-JAN-2004 ADE48853;

Human PRO 298 PCR primer #3.

Human; ss; PCR; secreted protein; transmembrane protein; PRO; cytostatic; ophthalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth; retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.

19-OCT-2001; 2001US-00166709

98WO-US021141. 98WO-US024855. 99WO-US000106.

06-JAN-2000; 2000WO-05000277. 06-JAN-2000; 2000WO-05000376. 11-FEB-2000; 2000WO-05000376. 12-FEB-2000; 2000WO-05006319. 22-FEB-2000; 2000WO-05006319. 10-MAR-2000; 2000WO-05006319. 17-MAR-2000; 2000WO-05006319. 17-MAR-2000; 2000WO-0501532. 30-MAR-2000; 2000WO-0501532. 30-MAR-2000; 2000WO-05014042. 30-MAR-2000; 2000WO-05015264. 31-MAR-2000; 2000WO-05015264. 32-MAR-2000; 2000WO-05015264. 32-MAR-2000; 2000WO-05015264. 32-MAR-2000; 2000WO-05015264. 32-MAR-2001; 2000WO-05019650. 32-MAR-2001; 2001WO-05019692. 30-JUL-2001; 2001WO-05019692. 30-JUL-2001; 2001WO-05019692. 2000WO-US00219. 2000WO-US00277. 2000WO-US00376. 2000WO-US034341. 2000WO-US004341. 99WO-US005190 99WO-US010733 99WO-US0128313 99WO-US028551 99WO-US028565 99WO-US031243 99WO-US031243 08-MAR-1999; 10-MAR-1999; 14-MAY-1999; 02-JUN-1999; 30-0NOV-1999; 02-DEC-1999; 16-DEC-1999; 16-DEC-1999; 30-DEC-1999; 05-JAN-2000;

(GETH) GENENTECH INC.

Ashkenazi AJ, Baker KP, Botetein D, Desnoyers L, Eaton DL; Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME; Goddard A, Goddwski PJ, Grimaldi JC, Gurney AL, Hillan KJ; Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL; Stewart TA, Tumas D, Williams PM, Wood WI;

WPI; 2004-008994/01.

New isolated nucleic acid encoding a PRO polypeptide, e.g. PRO4993 or PRO337, useful in molecular biology, chromosome and gene mapping, in generating antisense RNA and DNA, and in gene therapy.

Example 95; SEQ ID NO 519; 460pp; English.

The invention relates to an isolated PRO polypeptide (secreted or transmembrane protein) having at least 80% amino acid sequence identity transmembrane protein) having at least 80% amino acid sequence deneral control of including PRO lacking its associated signal peptide, a PRO extracellular domain with or without its associated signal peptide, a PRO extracellular domain with or without its associated signal peptide, a Also included are nucleic acids encoding the PRO proteins comprising the vector comprising a PRO nucleic acid), a host cell comprising the vector and producing PRO, a chimaeric molecule comprising to profused to a heterologous amino acid sequence, and an anti-PRO antibody. PRO337 polypeptide is useful for detecting a PRO4939 polypeptide is useful for detecting a proteptide. Similarly, PRO4939 polypeptide is useful for detecting profuse profuse and PRO739 polypeptide is useful for linking a bioactive molecule to a cell expressing PRO337 polypeptide is useful for linking a collective molecule to a cell expressing PRO4937 polypeptide; in the bioactive molecule to a cell expressing PRO4939 polypeptide; bioactive molecule coll. PRO337 polypeptide is useful for linking a causes death of the cell. PRO337 polypeptide is useful for linking a collecule to a cell expressing PRO4939 polypeptide; PRO725, PRO700 or PRO337 polypeptide is useful for linking a collecule to a cell expressing PRO4939 polypeptide; PRO725, PRO700 or PRO337 polypeptide is useful for linking a collecule to a cell expressing PRO4933 polypeptide; PRO725, PRO700 or PRO739 polypeptide is useful for linking a bioactive molecule to a cell expressing PRO4933 polypeptide; PRO725, PRO700 or PRO739 polypeptide is useful for linking a bioactive molecule to a cell expressing PRO4933 polypeptide; PRO7055, PRO700 or PRO739 polypeptide are useful for linking a bioactive molecule to a cell expressing PRO4933 polypeptide; PRO7055, PRO700 or PRO739 polypeptide are useful for linking a bioactive molecule to a cell expressing PRO4933 polypeptide; PRO7055,

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useful for linking a bioactive molecule to a cell expressing PRO155, useful for linking a bioactive molecule to a cell expressing PRO25, PRO700 or PRO739 polypeptide. PRO4993 polypeptide or anti-PRO37 polypeptide is useful for modulating at least one biological activity of the cell expressing PRO337 polypeptide, where the cell is killed. PRO337 polypeptide, where the cell is killed. PRO337 polypeptide is useful for modulating the biological activity of the cell expressing PRO493 polypeptide or an anti-PRO1559 polypeptide; PRO725, PRO700 or PRO739 polypeptide or an anti-PRO1559 polypeptide; BRO725, polypeptide, and PRO1559 polypeptide or anti-PRO725, anti-PRO706 or anti-PRO739 polypeptide is useful for modulating the biological activity of the cell expressing PRO725, PRO700 or PRO739 polypeptide are useful for modulating the biological activity of the cell expressing PRO725, PRO700 or PRO739 polypeptide are useful for modulating the biological activity of the cell expressing pro75, PRO700 or PRO739 polypeptide are useful for inhibiting tumour growth, retinal disorders, sports-related joint problems, articular cartilage defects, osteoarthritis or rheumatoid arthritis, wound healing and hearing loss in mammals. The present sequence is a PCR primer used to isolate nucleic acid encoding a PRO protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; ss; PCR; secreted protein; transmembrane protein; PRO; oytostatic; ophthalmological, antiarchritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth; retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                0; Gaps
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                                                                                                                                                                                                                                                                                                                                    Sequence 18 BP; 4 A; 6 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                               149 GGAGGCCGGCTTCGACT 165
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97US-00663411P-
97US-0066341P-
98US-0077450P-
98US-0077641P-
98US-0077641P-
98US-0077641P-
98US-0077641P-
98US-0077643P-
98US-0078910P-
98US-0078910P-
98US-0078918P-
98US-0078919P-
98US-007891P-
98US-007891P-
98US-0079664P-
98US-0079664P-
98US-0079664P-
98US-0079664P-
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ID ADE89954 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17 GGAGGTCGACTTCCACT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human PRO 298 PCR primer #3.
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10 - MAR - 1998
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12 - MAR - 1998
20 - MAR - 1998
25 - MAR - 1998
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27-MAR-1998;
27-MAR-1998;
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Matches
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PR 27-MAR-1998; 96US-0079276P.
PR 30-MAR-1998; 96US-0079922P.
PR 31-MAR-1998; 96US-0079922P.
PR 31-MAR-1998; 96US-0080165P.
PR 31-MAR-1998; 96US-0080165P.
PR 31-MAR-1998; 96US-0080137P.
PR 31-MAR-1998; 96US-0080137P.
PR 01-APR-1998; 96US-0081137P.
PR 15-APR-1998; 96US-008125P.
PR 15-APR-1998; 96US-008125P.
PR 22-APR-1998; 96US-008132P.
PR 22-APR-1998; 96US-00812P.
PR 22-APR-1998; 96US-00812P.
PR 22-APR-1998; 96US-00812P.
PR 22-APR-1998; 96US-0081
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2000WO-US000219-
2000WO-US000217-
2000WO-US0003565-
2000WO-US00036319-
2000WO-US005819-
2000WO-US005819-
2000WO-US005819-
2000WO-US005819-
2000WO-US005819-
2000WO-US005819-
2000WO-US005819-
2000WO-US014941-
2000WO-US01494-
2001WO-US01494-
2001WO-US01494-
2001WO-US01494-
2001WO-US01494-
98US-0094651P

98US-010038P

98US-0100334P

98UG-0103248E5

98UG-0113228E5

99UG-013621P

99UG-013621P

99UG-013621P

99UG-013621P

99UG-013622P

99UG-013623P

99UG-013628P

99UG-013628P

99UG-013628P

99UG-0146588P

99UG-0146588P

99UG-014658P

99UG-014658P
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2001WO-US021735
2001US-00918585
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GERBER H.
GERRITSEN M E.
GODDARR A.
GODOWSKI P J.
GODOWSKI P J.
GIRNBEY A L.
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BOTSTEIN D.
DESNOYERS L.
EATON D L.
FERRARA N.
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                                                                        20-NOV-1998;
22-DEC-1998;
23-DEC-1998;
05-JAN-1999;
08-MAR-1999;
10-MAR-1999;
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28-JUL-1999;
29-OCT-1999;
30-NOV-1999;
02-DEC-1999;
                                                                                                                                                                                         12-MAR-1999;
29-MAR-1999;
21-APR-1999;
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14-MAY-1999;
02-JUN-1999;
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(BAKE/)
(BOTS/)
(DESN/)
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(FILV/)
(FONG/)
(GAOW/)
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(GODD/)
(GODO/)
(GIRM/)
(GURN/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the N-terminal and lysine residues.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group; bioavailibility; formulation; neurological disorder; inflammatory disorder; autoimmune disorder; cancer; neurodegenerative disorder; parkinson; disease; Huntington's disease; Alzheimer's disease; neurological injury; stroke; multiple sclerosis; malignant glioma; medulloblastoma; neuroectodermal tumour; ss.
                                                                                                                                                                                                                           °,
                                                                                                                                                                                          Query Match

2.9%; Score 12.2; DB 1; Length 18;
Best Local Similarity 82.4%; Pred. No. 4.7e+02;
Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:45.
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                                                                                                                                                                                                                                                       149 GGAGGCCGGCTTCGACT 165
                                                                                                                                                                                                                                                                                                                                                            AAF27041 standard; DNA; 35 BP.
                                                                                                                                                                                                                                                                         17 GGAGGTCGACTTCCACT 1
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13-AUG-1999; 99US-0149016P.
                                                                                                                                                                                                                                                                                                                                                                                                                         30-MAR-2001 (first entry)
                                       PAN J.
PAN J.
ROY N F.
SHELTON D L.
STEWART T A.
TUMAS D.
WILLIAMS P M.
KLJAVIN I J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-049927/06.
             KUO S S.
NAPIER M A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
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                                                                                                  (STEW/)
(TUMA/)
(WILL/)
(WOOD/)
             (KUOS/)
(NAP1/)
(PANJ/)
(PAON/)
(ROYM/)
KLJA/)
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defining and mapping functionally important regions of a protein by modifying accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of various medical conditions including various neurological disorders.

Inflammatory and autofimmune diseases, and cancers. In particular, they may be used to prevent preventing or amaliorate neurodegenerative disorders (e.g., Parkinson's disease, and cancers. In particular, they may be used to prevent preventing or amaliorate neurodegenerative disorders (e.g., Parkinson's disease, neurological injury and trauma; immunological diseases, the nervous system (e.g., multiple sclerosis); stroke; and malignant gliomas, medulloblastomas and trauma; immunological diseases of the nervous system (e.g., multiple sclerosis); stroke; and malignant gliomas, medulloblastomas and result in increased half-life, altered tissue distribution (such as an improved ability to stay in the vasculature for longer periods of time); increased stability in solution, protection from proteolytic degradation, or reduced immunogenicity. In particular, the ability to remain in the vasculature for prolonged periods may allow a hedgehog protein of the invention to cross the blood-brain barrier, and an increased thermal stability would be an advantage when formulating the hedgehog protein in powder form. The present sequence represents a human invention of the manning the headen of the manning the headen of the headen 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.9%; Score 12.2; DB 1; Length 35; 68.0%; Pred. No. 1.1e+03; ative 0; Mismatches 8; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 35 BP; 8 A; 15 C; 9 G; 3 T; 0 U; 0 Other;
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920S-00882713.
920S-00882714.
920S-00882824.
920S-00882886.
920S-00882886.
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(first entry)
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14-MAY-1992;
14-MAY-1992;
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14-MAY-1992;
14-MAY-1992;
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26-MAY-1994
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14-MAY-1992;
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92US-00882922. 92US-00883823. 92US-00883849.

14-MAY-1992

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The sequences (AAQ52923-Q53037) are pref. herpes simplex virus target sequences for enzymatic RNA molecules. The RNA molecules are complementary to a substrate binding region in the specified gene target. They also have enzymatic activity, in that they specifically cleave RNA in the target. The ERMs interfere with viral replication and therefore have anti-viral properties. They can be used to attenuate viruses to be used in vaccines. (Updated on 25-MAR-2003 to correct RN field.) (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                       Enzymatic RNA molecules - used to inhibit viral replication, infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tag sequence; colorectal cancer; pancreatic cancer; colon cancer; diagnosis; prognosis; treatment; ss.
                                                                                                                                                                                                                          Dudycz LW, Mcswiggen JA, Macejak DG, Holecek JJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tag sequence of a transcript increased in pancreatic cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.8%; Score 12; DB 1; Length 13; 91.7%; Pred. No. 2.6e+02; Live 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 13 BP; 4 A; 3 C; 5 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                              Claim 5; Fig 15; 287pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX31516 standard; DNA; 15 BP.
                    9205-00884433
9205-00884432
9205-00884431
9205-00884436
9205-00884521
9205-00923738
9205-00935864
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                                                                                                                                     92US-00963322
92US-00987129
                                                                                                                                                             92US-00987130
                                                                                                                                                                                                   (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (UYJO) UNIV JOHNS HOPKINS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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Best Local Similarity 91.7
Matches 11, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 cuccaccadada 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        32 CTGGGACGAAGA 43
                                                                                                                                                                                                                                                               WPI; 1993-386599/48.
                                                                                                                                                                                                                                                                                                     and gene expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-MAY-1997;
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                                                                                                                                                                          07-DEC-1992;
                                                                                                26-AUG-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-NOV-1998.
                                                                                                                        18-SEP-1992;
                                                                                                                                                                                                                           Draper KG,
Mamone JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX31516;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 906
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Use of isolated gene transcripts - useful for developing products for the diagnosis, prognosis and treatment of cancers, particularly colon and pancreatic cancer. Claim 13; Page 57; 120pp; English Vogelstein B, Kinzler KW; WPI; 1999-070161/06

differentially expresent tag sequences of transcripts that are differentially expressed in colorectal cancer, in pancreatic cancer, or in both. The tag sequences can be used to identify genes by matching the tag to a gen data base member, or by using the tag sequences as probes to isolate unidentified genes from cDNA libraries. The tag sequences can slow be used in a method for diagnosing colon or pancreatic cancer in a sample suspected of being neoplastic. The method comparises comparing the level of at least one transcript in a first sample of a tissue to a second sample, where the first sample is a colonic tissue suspected of being neoplastic and the second sample is a normal human colonic tissue. The transcript is identified by a tag selected from AAX30947-1815. The methods of the invention can be used in the diagnosis, prognosis and treatment of cancer

Sequence 15 BP; 3 A; 2 C; 6 G; 4 T; 0 U; 0 Other;

0; Gaps Query Match 2.8%; Score 12; DB 1; Length 15; Best Local Similarity 100.0%; Pred. No. 3.5e+02; Matches 12; Conservative 0; Mismatches 0; Indels

92 CATCACCACGIC 103

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AAF48829 standard; DNA; 15 AAF48829; AAF48829

BP.

IGFBP3 oligonucleotide #2249. 30-MAR-2001 (first entry)

Antisense therapy, antiproliferative, antinflammatory, antipsoriatic, cytostatic, dermatological; cardiant, virucide, ophthalmological; keloid; skin discorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding procein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keardosis; neophasia; scleroderme, wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplama; kidney disease; neovascular condition; hyperplama; kidney disease;

Homo sapiens.

WO200078341-A1.

21-JUN-2000; 2000WO-AU000693. 28-DEC-2000

21-JUN-1999;

Werther GA, Edmondson SR; Wraight CJ,

MURD-) MURDOCH CHILDRENS RES INST.

WPI; 2001-041421/05

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that

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The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisomediac procein (for Insulin-like Growth Factor [IGP]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBBJ), which is capable of inhibiting or reducing growth factor mediated cell proliferation, infilammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF4151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrheea, keloids, keratosis, chthyosis, galeroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic condition as the properpoliferation of the inside of blood
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
reduces growth factor mediated cell proliferation and/or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
2.8%; Score 12; DB 1; Length 15;
Best Local Similarity 100.0%; Pred. No. 3.5e+02;
Matches 12; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15 BP; 6 A; 4 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                Example 7; Page 58; 201pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    vessels or any other hyperplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  394 CCAAGAAGGTCT 405
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     inhibits or r
inflammation.
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Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin discorder; insulin-like Growth Factor I receptor; IGFP-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilatis; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scaleroderma; wart; skin cancer; sclerotic disease; hypermeovascular condition; hyperplasia; kidney disease; neobascular condition; hyperplasia; kidney disease; IGFBP3 oligonucleotide #2252.

AAF48832 standard; DNA; 15 BP.

RESULT 908 AAF48832

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30-MAR-2001 (first entry)

AAF48832;

Homo sapiens.

40200078341-A1.

21-JUN-2000; 2000WO-AU000693.

99US-0140345P. 21-JUN-1999; (MURD-) MURDOCH CHILDRENS RES INST

Wraight CJ, Werther GA, Edmondson SR;

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.

Example 7; Page 58; 201pp; English

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an anticontection (for Insulin-like Growth Pactor [1678]—1 receptor, IGF binding protein [1678]—2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonuclectide which can be used to design the artisense oligonuclectide which can be used to design the artisense oligonuclectide of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, chthyosis, pityriasis, ruba, pilaris, serborthoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hypernecvascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies other sclerotic disease, kidney disease, hyperproliferation of the inside of blood to sessels or any other hyperplasia

Sequence 15 BP; 4 A; 4 C; 5 G; 2 T; 0 U; 0 Other;

Gaps ö / Match 2.8%; Score 12; DB 1; Length 15; Local Similarity 100.0%; Pred. No. 3.5e+02; Local 12; Conservative 0; Mismatches 0; Indels Query Match Best Loca Matches

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AAF48830 standard; DNA; 15 RESULT 909 AAF48830

AAF48830;

30-MAR-2001 (first entry)

IGFBP3 oligonucleotide #2250.

Antisense therapy, antiproliferative, antiinflammatory, antipsoriatic, cytostatic, dermatological, cardiant; virucide, ophthalmological, keloid, sein discorder, insulin-like Growth Factor. I receptor; IGF-1; pityriasis; IGF binding protein; IGFB-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; scrborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neobascular condition; hyperplasia; kidney disease;

Ното варіеля

WO200078341-A1

21-JUN-2000; 2000WO-AU000693.

99US-0140345P 21-JUN-1999;

(MURD-) MURDOCH CHILDRENS RES INST.

Wraight CJ, Werther GA, Edmondson SR;

WPI; 2001-041421/05

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.

Example 7; Page 58; 201pp; English.

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1

; 0 receptor, IGF binding protein (IGFBP)-2 or IGFBPB), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antiesnse oligonucleotides of the present invention (see AAF45151 and AAF45153-F55161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplastas, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperpoliferation of the inside of blood vessels or any other hyperplasia Gaps ., Query Match
2.8%; Score 12; DB 1; Length 15;
Best Local Similarity 100.0%; Pred. No. 3.5e+02;
Matches 12; Conservative 0; Mismatches 0; Indels Sequence 15 BP; 5 A; 4 C; 4 G; 2 T; 0 U; 0 Other; 394 CCAAGAAGGTCT 405 3 CCAAGAAGGTCT 14 8888888888888888 ਨੋ 셤

AAF48831 standard; DNA; 15 BP. RESULT 910

(first entry) 30-MAR-2001 AAF48831;

IGFBP3 oligonucleotide #2251.

Antisense therapy, antiproliferative, antiinflammatory, antipsoriatic, cytostatic, dermatological, cardiant, virucide, ophthalmological, keloid, stin disorder, Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis, IGF binding protein, IGFBP-2, IGFBP3, inflammation, psoriasis, pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea, ruba; keratosis; neoplasia; sclaroderma; wart, skin cancer; sclerotic disease; hypermedvascular condition; hyperplasia, kidney disease; neoblar condition of the retina; ss.

Homo sapiens.

WO200078341-A1.

21-JUN-2000; 2000WO-AU000693.

21-JUN-1999; 99US-0140345P.

(MURD-) MURDOCH CHILDRENS RES INST.

Wraight CJ, Werther GA, Edmondson SR;

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.

Example 7; Page 58; 201pp; English.

ij The present invention relates to a method for ameliorating the effects skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein (IGFBP]-2 or IGFBPP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense

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Page 485
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oligonuclectides of the present invention (see AAF45151 and AAF45153-
P45161). The method is useful for ameliorating the effects of psoriasis,
inchthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis,
neoplasias, scleroderma, warts, benign growths, cancers of the skin, a
hyperneovascular condition such as a neovascular condition of the retina,
brain or skin, growth factor-mediated malignancies, other sclerotic
disease, kidney disease, hyperpoliferation of the inside of blood
vessels or any other hyperplasia
               888888888888888
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Sequence 15 BP; 4 A; 4 C; 5 G; 2 T; 0 U; 0 Other;

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0
                                           Gaps
                                         .
/ Match 2.8%; Score 12; DB 1; Length 15; Local Similarity 100.0%; Pred. No. 3.5e+02; les 12; Conservative 0; Mismatches 0; Indels
         Query Match
                                               Matches
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394 CCAAGAAGGICT 405 CCAAGGAGGTCT 13 ઠે

AAS99932 standard; DNA; 15 BP. 12-MAR-2002 AAS99932; RESULT 911 AAS99932

Bven-skipped homeobox 1 (EVX1) gene allele-specific oligonucleotide #9 (first entry)

Even-skipped homeo box 1; EVX1; neurological disease; drug screening; haplotyping; single nucleotide polymorphism; SNP; human; ss; allele-specific oligonucleotide.

domo sapiens.

MO200190120-A2.

29-NOV-2001

21-MAY-2001; 2001WO-US016559.

19-MAY-2000; 2000US-0205437P.

(GENA-) GENAISSANCE PHARM INC

Kumar AM; Kliem SE, Duda A,

WPI; 2002-089913/12.

Novel genetic variants of even-skipped homeo box 1, EVX1 gene useful for therapeutic purposes and for expressing EVX1 protein useful in identifying drugs to treat neurological diseases.

Claim 16; Page 13; 69pp; English.

The invention relates to an isolated polynucleotide (I), comprising a nucleotide sequence which is a polymorphic variant of a reference sequence for the even-skipped homeo box 1 (homologue of Drosophila) sequence for the even-skipped homeo box 1 (homologue of Drosophila) cutto. EUXI) gene or its fragment, or a polymorphic variant of a reference sequence for a EVXI cDNA or its fragment. EVXI polypeptide (II) is useful for sercening for drugs targeting the polypeptide, by contacting the EVXI polymorphic variant with a candidate agent and assaying for binding colymorphic variant with a candidate agent and assaying for binding activity. A method is described for identifying an association between a trait such as a clinical response to a drug targeting EVXI and a haplotype or haplotype pair of EVXI gene. The methods are useful in developing diagnostic tests and therapeutic treatments for neurological diseases. (I) is useful for studying the expression and function of EVXI and expressing EVXI protein for use in screening for candidate drugs to treat diseases related to EVXI activity. The polymorphism and haplotype cut duras to treat neurological diseases, screening for euch drugs and reducing bias in clinical trials of such drugs. (I) is useful for

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therapeutic purposes. (I) is useful for determining if an individual has one of the haplotypes 1.4 or the haplotype pairs. Establishing the EVX1 one of the haplotype pair of an individual is useful for improving the efficiency and reliability of several steps in the discovery and considered to the expension of the efficiency and reliability of several steps in the discovery and control of a newlogical diseases. The haplotyping method is useful to validate EVX1 as a candidate target for treating a specific condition or disease content of the EVX1 segences in vivo screening and testing of drugs against EVX1 protein and for testing the efficacy of therapeutic agants and compounds for neurological diseases in a contingual diseases in a prognostic formats and therapeutic methods, for immunoprecipitating condition, for detecting EVX1 protein isoforms in biological condition, for detecting EVX1 protein isoforms in biological conditions, calls which have been fixed or unfixed and prepared on slides, for use in immunocytochemical, immunohistochemical and immunofluorescence techniques. AAS99958 immunication in the effect of the expension of the efficiency of the immunofluorescence techniques.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15 BP; 2 A; 6 C; 6 G; 0 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention
                       $$$$$$$$$$$$$$$$$$$$$$$
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2.8%; Score 12; DB 1; Length 15; 85.7%; Pred. No. 3.5e+02; iive 1; Mismatches 1; Indels 2.007 Best Local Similarity 85.79 Matches 12, Conservative

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Gaps ô

> ABK70537 standard; DNA; 15 BP (first entry) 15-JUL-2002 ABK70537; RESULT 912

Human G protein-coupled receptor 7 allele-specific probe #21. Human; G protein-coupled receptor 7; GPR7; haplotyping; SNP; psychological disorder; neurological disorder; probe; 88; single nucleotide polymorphism.

Homo sapiens.

WO200222644-A1.

21-MAR-2002.

17-SEP-2001; 2001WO-US029207.

(GENA-) GENAISSANCE PHARM INC. 15-SEP-2000; 2000US-0232900P.

Sanchis A, Koshy B,

WPI; 2002-383121/41.

Novel genetic variants of G protein-coupled receptor 7 gene useful for therapeutic purposes and for expressing GPR7 protein useful in identifying drugs to treat psychological and neurological disorders.

Claim 16; Page 13; 69pp; English.

The invention relates to an isolated polynuclectide (I) comprising a nucleotide sequence which is a polymorphic variant of a reference sequence for G-protein coupled receptor 7 (GPR7) gene or its fragment, or a polymorphic variant of a reference sequence for a GPR7 cDNA or its fragment. The encoded polypeptide (II) is useful for screening for drugs targeting the polypeptide. (I) is useful for identifying an association

cc between a trait such as a clinical response to a drug targeting GPR7 and a applictype or haplotype pair of GPR7 gene. Such methods have applictability in developing diagnostic tests and therapeutic treatments or applictability in developing diagnostic tests and therapeutic treatments payliciability in developing diagnostic tests and therapeutic treatments of contrological and neurological disorders. (I) is useful for validating whether contrological disorders, screening to treat psychological and reducing bias in claimable target for drugs to treat psychological and neurological disorders, screening for such drugs and reducing bias in claimable target for contrological disorders, screening for such drugs and reducing bias in claimability the GRP7 haplotype pair of an individual is useful for improving the efficiency and reliability of several steps in the discovery and development of drugs for treating diseases associated with depty and sevent of drugs for treating diseases associated with GRP7 activity psychological and neurological disorders. The haplotype method is useful to validate GRP7 activity. The method is useful to validate GRP7 activity. The method is also useful in screening for compounds to the drugs and neurological disorders as sociated with GRP7 activity psychological and neurological disorders or to population and the method is a specific condition or disease predicted to be associated with GRP7 activity or condition or disease predicted to be associated with GRP7 activity or care in the method is the most frequent GRP7 isoforms present in the compounds that display the highest desired agonist or antagonist activity for each of the most frequent GRP7 isoforms present in the most frequent GRP7 isoforms and neurological disorders and in assays to measure the binding affinities of one or more candidate drugs targeting GRP7 for the treatment of psychological and neurol

Sequence 15 BP; 3 A; 4 C; 7 G; 0 T; 0 U; 1 Other;

Gaps . 0 2.8%; Score 12; DB 1; Length 15; llarity 85.7%; Pred. No. 3.5e+02; Conservative 1; Mismatches 1; Indels Best Local Similarity Matches 12; Conserv Query Match

333 GACGACCAGGCCG 346

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caccadecacecse 15 원

ABN80596 standard; DNA; 15 BP ABN80596; RESULT 913

19-JUL-2002 (first entry)

Human P450(cytochrome) oxidoreductase allele specific PCR primer #36.

P450 (cytochrome) oxidoreductase; POR; cancer; haplotype; SNP; nuclectide polymorphism; flavoprotein; enzyme; PCR; primer; ss Human; single

Homo sapiens

WO200226768-A2

04-APR-2002

01-OCT-2001; 2001WO-US030877

(GENA-) GENAISSANCE PHARM INC. 29-SEP-2000; 2000US-0236449P. Kazemi A, Kliem SE,

Tanguay DA; Lanz EM, Messer C,

WPI; 2002-394236/42.

New genetic variants comprising haplotypes of the P450 (cytochrome) oxidoreductase (POR) isogene, useful in improving the efficiency of drug screening protocols for compounds targeting POR. Claim 14; Page 15; 141pp; English. The present invention provides the protein, gene and cDNA sequences of human P450(cytochrome) oxidoreductase POR, and single nucleotide polymorphisms (SNPs) identified therein. The sequences can be used to haplotype the POR gene of an individual, and to establish whether POR is a suitable target for drugs to treat cancer and disorders associated with impaired protein synthesis in cells. The present sequence is an allele specific primer for the coding sequences of the invention

Sequence 15 BP; 2 A; 9 C; 2 G; 1 T; 0 U; 1 Other;

Gaps ; 0 Score 12; DB 1; Length 15; Pred. No. 3.5e+02; 1; Mismatches 1; Indels Query Match
Best Local Similarity 85.7%;
Matches 12; Conservative

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ABN87913 standard; DNA; 15 BP. RESULT 914

ABN87913;

12-AUG-2002 (first entry)

Human GSR allele specific oligonucleotide primer SEQ ID NO:32.

Human, glutathione reductase; GSR, enzyme; haemolytic anaemia; SNP; gene therapy; antianaemic; polymorphic; single nucleotide polymorphism; orimer; BB

Homo sapiens

/*tag= a /note= "polymorphic base" Location/Qualifiers misc_feature

WO200242320-A2 30-MAY-2002

13-NOV-2001; 2001WO-US046473.

10-NOV-2000; 2000US-0247202P.

Sausker EA, (GENA-) GENAISSANCE PHARM INC.

Bieglecki KW, Sanchis A,

WPI; 2002-471719/50.

New genetic variants of Glutathione reductase isogenes, useful for improving efficiency and reliability in drug development for treating hemolytic anemia.

Claim 14; Page 14; 137pp; English

The present invention describes genetic variants of the human glutathione reductase (GSR) gene (I). (I) has antianaemic activity and can be used in gene therapy. (I) can be used in screening for drugs targeting (I) that are useful for treating haemolytic anaemia. Methods from the present invention can be used: for improving the efficiency and reliability of several steps in the discovery and development of drugs for treating diseases associated with GSR activity; for haplotyping, which is also used by the pharmaceutical research scientist to validate GSR as a candidate target for treating a specific condition of disease predicted to be associated with GSR activity, e.g. haemolytic anaemia, and in the case associated with GSR activity, and for screening compounds targeting GSR. (I) is useful in studying the expression and function of GSR, and in expressing GSR protein for use in screening compounds targeting GSR. (I) is useful in studying the expression and function of GSR, and in expressing affinity of candidate drugs to creating GSR for the treatment of haemolytic anaemia. The present sequence represents an allele specific of haemolytic anaemia. The present sequence represents an allele specific condition of the present invention. N.B. The polymorphic base (showing a single nucleotide polymorphism) in the ASO primer is shown cusing an IUPAC ambiguity code (as given in the present invention)

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Sequence 15 BP; 1 A; 8 C; 4 G; 1 T; 0 U; 1 Other;

ö Gaps ô 2.8%; Score 12; DB 1; Length 15; 85.7%; Pred. No. 3.5e+02; tive 1; Mismatches 1; Indels Local Similarity 85.7 nes 12; Conservative Query Match Best Lcc Matches

320 CGTGCTGGCGGCGG 333

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RESULT 915

ABL51980 standard; DNA; 15 BP ABL51980

(first entry) 11-JUL-2002 ABL51980;

Human SLC18A2 allele specific oligonucleotide primer SEQ ID NO:28.

Human; solute carrier family 18 member 2; SLC18A2; vesicular monoamine; vesicular monoamine transporter; VMAT2; polymorphic site; SNP; single mucleotide polymorphism; antiinflammatory; neuroleptic; haplotyping; genotyping; respiratory inflammatory disease; neuropsychiatric disorder; monoaminergic brain system; primer; ss.

Homo sapiens

Location/Qualifiers misc_feature

/*tag= a /note= "polymorphic site indicated by an ambiguity base"

WO200222652-A2

21-MAR-2002

.7-SEP-2001; 2001WO-US042217

15-SEP-2000; 2000US-0232895P

(GENA-) GENAISSANCE PHARM INC

Sausker EA; Kliem SE, Anastasio AE,

WPI; 2002-393942/42.

Novel genetic variants of soluble carrier family 18 (vesicular

monoamine), member 2 gene useful for screening drugs to treat diseases e.g. neuropsychiatric disorders involving monoaminergic brain systems.

Claim 17; Page 14; 183pp; English.

The present invention describes an isolated polynuclectide (I) having a sequence (SI) comprising soluble carrier family 18 (vesicular monoamine), member 2 (SLC18A2) isogene selected from 49 isogenes with regions of a sequence (SS) of 40023 bp (see ABL51954), and defined by a corresponding set of polymorphisms whose locations and identities are given in the specification; or a sequence (S2) complementary to (S1). (I) has continuismantary to (S1). (I) has continuismantary to set of polymorphisms whose locations and invention can be used for haplotyping the SLC18A2 gene in an individual. SLC18A2 is also known and genotyping the SLC18A2 gene in an individual. SLC18A2 is also known as the vesticular monoamine transporter (WMAT2). (I) is useful in studying the expression and function of SLC18A2, and in expressing the SLC18A2 continuity of the biological activity of SLC18A2, and in expressing the surfation conthe biological activity of SLC18A2 as well as on the biological activity of SLC18A2 is well as on the binding affinity of candidate drugs targeting SLC18A2 for the treatment of respiratory conthe biological activity of SLC18A2 is an electrof slowders involving concommency brain systems. The present sequence represents an allele specific oligonuclectide (ASO) primer for human SLC18A2, which is given in the present invention

Sequence 15 BP; 2 A; 7 C; 5 G; 0 T; 0 U; 1 Other;

Gaps ; Query Match

2.8%; Score 12; DB 1; Length 15;
Best Local Similarity 85.7%; Pred. No. 3.5e+02;
Matches 12; Conservative 1; Mismatches 1; Indels

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106 ACCGCGACCGCAGC 119

ઠે a RESULT 916

AAS19726/c ID AAS19726 Btandard; DNA; 15 BP.

AAS19726;

08-MAY-2002 (first entry)

ASO probe #23 to detect human RANGAP1 gene polymorphisms.

Human; single nucleotide polymorphism; SNP; RANGAP1; haplotyping chromosome 22q13.2-q13.31; Ran GTPase activating protein 1; genotyping; cancer; irregular cell cycle associated disorder; ASO; probe; ss; allele-specific oligonucleotide.

Homo sapiens

40200179240-A2.

25-OCT-2001

17-APR-2001; 2001WO-US012455.

17-APR-2000; 2000US-0198072P.

(GENA-) GENAISSANCE PHARM INC.

Choi JY, Koshy B; Chew A,

WPI; 2002-075068/10.

Genotyping human Ran GTPase activating protein 1 gene of individual for determining haplotype of individual, involves determining identity of nucleotide pair at specific polymorphic sites for two copies of the gene.

Claim 15; Page 14; 148pp; English

Page 488

The present invention relates to novel single nucleotide polymorphisms (SNPs) in the human Ran GTPase activating protein 1 (RANGAP1) gene located on chromosome 20413.2-413.31, and methods for haplotyping and/or genotyping the RANGAP1 gene. The methods of the invention make use of allele-specific oligonucleotides (ASOs) as probes and primers and/or polymorphisms. The polymucleotides for detecting the RANGAP1 gene polymorphisms. The polymucleotides and screened compounds are useful for treatment of diseases associated with RANGAP1 activity, such as cancer and other disorders associated with an irregular cell cycle. AAS19704-AAS19742 represent ASO probes for detecting human RANGAP1 gene

886666666666688888

Sequence 15 BP; 2 A; 5 C; 4 G; 3 T; 0 U; 1 Other;

Gaps .. 0 2.8%; Score 12; DB 1; Length 15; 85.7%; Pred. No. 3.5e+02; iive 1; Mismatches 1; Indels 56 AGAGGAGTCTCTGC 69 Local Similarity 85.7 nes 12; Conservative Query Match Best Loc Matches ઠે

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15 AGAGGAGYCCCTGC 2 g

AAS97315 standard; DNA; 15 AAS97315; RESULT 917 AAS97315/c

BP.

(first entry) 12-MAR-2002 Human CRYBB1 gene ASO probe #10.

Human, crystallin beta B1; CRYBB1; chromosome 22q12.1; ophthalmalogical; cataract; allele specific oligonucleotide; ASO; probe; ss; haplotype; genotyping; transgenic animal.

Homo sapiens.

WO200185998-A1.

15-NOV-2001

07-MAY-2001; 2001WO-US014715.

05-MAY-2000; 2000US-020253P.

(GENA-) GENAISSANCE PHARM INC

Choi JY, Kazemi A, Kliem SE,

Koshy B, Rounds E;

WPI; 2002-062253/08.

Novel polymorphic variants of crystallin, beta B1 useful in studying expression and function of the protein, useful for screening candidate drugs to treat diseases e.g. cataract.

Claim 15; Page 12; 94pp; English.

The invention relates to an isolated polynucleotide comprising a sequence which is a polymorphic variant of a reference sequence for crystallin, where the (GYRBH). located on chromosome 22012.1) gene or their fragment, where the polymorphic variant comprises a CRYBH isogene defined by a haplotype from haplotypes 1-16 as given in the specification. Also included are a transgenic non-human animal transformed or transfected with the polymorphic variant, a computer system for storing and analysing polymorphism data for CRYBH gene, a genome anthology for the CRYBH gene which comprises the defined CRYBH isogenes, methods of determining an individuals haplotype or genotype as well as methods of determining the association of a particular haplotype with a disease or trait and a composition comprising at least one genotyping or trait and a (especially allele-specific oligonucleotides (ASO)) for detecting a polymorphism in the CRYBH. The isogenes or haplotypes are useful for

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improving the efficiency and reliability of several steps in the discovery and development of drugs for treating diseases associated with CRYBB1 activity, e.g. caearact, and can also be used by the pharmaceutical research scientist to validate CRYBB1 as a candidate traget for, and in design of clinical trials of candidate drugs for, treating a specific condition drugs or disease predicted to be associated with CRYBB1 activity. The ASOs are useful as probes and primers, and for assaying a polymorphism in the target region. The present sequence is an ASO probe for CRYBB1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New Pro-Platelet Basic Protein (PPBP) gene polymorphic variants, useful for studying the expression and function of PPBP and screening candidate drugs for treating disorders associated with PPBP activity, e.g. immunological disorders.
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0
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                                                                                                                                                                                 Query Match
2.8%; Score 12; DB 1; Length 15;
Best Local Similarity 85.7%; Pred. No. 3.5e+02;
Matches 12; Conservative 1; Mismatches 1; Indels
                                                                                                                                                      Sequence 15 BP; 3 A; 3 C; 8 G; 0 T; 0 U; 1 Other;
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                                                                                                                                                                                                                                                 132 CTGGCCCGCCTGGC 145
                                                                                                                                                                                                                                                                                                                                                         AAL46088 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                   15 CTGCCCCRCCTGGC 2
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ID AAL4
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Length 15; Sequence 15 BP; 2 A; 4 C; 5 G; 3 T; 0 U; 1 Other; Score 12; DB 1; Pred. No. 3.5e+02; 2.8%; Query Match Best Local Similarity

The present invention provides the protein, cDNA and genomic sequences of human pro-platelet basic protein (PPBP) and single nucleotide polymorphisms (SNPs) identified therein. The polymorphic variants are useful in studying the expression and function of PPBP, in expressing PPBP protein for use in screening for candidate drugs to treat diseases related to PPBP activity, in studying the effect of the variation on the biological activity of PPBP, and the binding affinity of candidate drugs targeting PPBP for the treatment of disorders associated with PPBP activity, e.g. metabolic and immunological disorders. The present sequence is an allele specific probe for the gene of the invention

Claim 14; Page 12; 68pp; English.

ABK32470

RESULT 919

ABK32470,

Matches

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New oligo nucleoside(s) and nucleotide(s) with up to 200 bases - nuclease resistant anti sense cpds. useful for treating hereditary disorders of altered genetic expression mechanisms.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Maize; corn; Zea mays; delta-9 desaturase; GBSS; target; substrate; granule bound starch synthase; hammerhead riboxyme; haltpin riboxyme; modulation; gene expression; transgenic plant; cleavage; canola plant; caffeine synthesis; coffee plant; nicotine production; tobacco; fruit ripening; flower pigmentation; lignin production; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Delta-9 desaturase hamerhead ribozyme target SEQ ID NO:829.
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2.8%; Score 12; DB 1; Length 16;
Best Local Similarity 100.0%; Pred. No. 4e+02;
Matches 12; Conservative 0; Mismatches 0; Indels
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                                                                             /*tag= a
/mod_base= OTHER
/note= "see comments"
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/mod_base= OTHER
/note= "see comments"
                                              location/Qualifiers
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90US-00582287.
90US-00582456.
90US-00582457.
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13-SEP-1990;
13-SEP-1990;
09-APR-1991;
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Moskwa PS,
                Synthetic
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AAX62954/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tetraethylene glycol; cancer; antisense; gene expression; inhibition;
diol; ss.
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 Gaps
                                                                                                                                                                                                                                                                                              Human, colon cancer, colorectal cancer, pancreatic cancer; SAGE tag, serial analysis of gene expression, diagnostic, prognostic, probe,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New human nucleic acid containing specific SAGE tags, useful as diagnostic markers for cancer, also derived probes.
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   Indels
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   Mismatches
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AAQ21895
ID AAQ21895 standard; DNA; 16 BP.
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                                                                                                                                                           ABK32470 standard; DNA; 15
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   12; Conservative
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                                    282 GGCACCAAGCTGGT
                                                       GGCACCARGCTAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-153821/20.
                                                                                                                                                                                                                                                                                                                                   cancer marker; ss
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Gaps

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13-JUL-1995;

WO9710328-A2 20-MAR-1997

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A method has been developed for the identification of a nucleic acid capable of modulating a process in a biological system. The method capable of introducing into the system a random library of nucleic comparises: (a) introducing into the system a random library of nucleic acid catalytic Garalytic domain (ED); and (b) identifying NPC or a random sequence, and a catalytic domain (ED); and (b) identifying NRC in systems where modulation has occurred and/or determining the sequence of at least part of the SBDs in such systems. Nucleic acid molecules with endonuclease activity and catalytic activity, from the present invention, are used to modulate gene expression in plant and mammalian calls and to cleave target nucleic acid, particularly for treating systemic diseases caused by specific NNA, e.g. cancer, inflammation, psoriasis, non-hepatic activity that modulate expression of the Raf gene, are used to treat cancer, references, psoriasis or rheumatoid arthritis, or used to treat cancer, references, psoriasis or rheumatoid arthritis, or used to treat cancer, references, with the level of c-raf. Introduction of sugar/phosphate modifications increases stability against nuclease and activity Anyslozz to AAV932Y7 represent NRS that contain the and activity and and activity represent name of the craft introduction activity and and activity represent name of activity and activity and activity represent name activity and activity and activity and activity the level of c-raf. Introduction of sugar/phosphate modifications increases stability against nuclease and activity and activity represent NRS that contains and activity an
                                                                                                                                                                                                                                                                                                                                                                                               Identifying new catalytic nucleic acid that modulates selected processes - especially ribozymes that cleave Raf RNA for treating cancer, restenosis, and also new ribozymes and modified nucleoside triphosphates used as antiviral agents and synthons.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         activity. AAV90922 to AAV93877 represent NACs that can be used in
method, specifically for modulating the expression of a Raf gene
                                                                                                                                                                                                                   Jarvis T, Matulic-Adamic J, Reynolds M, Kisich K, Bellon L.
Parry T, Beigelman L, Mcswiggen JA, Karpeisky A, Burgin A;
Thompson J, Workman CT, Beaudry A, Sweedler D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.8%; Score 12; DB 1; Length 17;
100.0%; Pred. No. 4.6e+02;
lve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 177; Page 157; 259pp; English.
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                        97US-0061321P.
97US-0061324P.
97US-0064866P.
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                                                                                                                97US-0068212P
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                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           396 AAGAAGGICTIC 407
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 2.8
Best Local Similarity 100.
Matches 12, Conservative
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22-AUG-1997;
02-OCT-1997;
02-OCT-1997;
05-NOV-1997;
                                                                                                                                                                                                                                                      Parry T, Be
Thompson J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABN01021;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes an enzymatic nucleic acid molecule (I) with RNA cleaving activity, which modulates the expression of a plant gene. Also described is a gene comprising a CNNA sequence encoding maize Delta-9 desaturase. (I) can be used to modulate expression of a gene, preferably Delta-9 desaturase or a granule bound starch synthase (GBSS) gene, in a plant (preferably a maize or canola plant). (I) can be used to modulate eaffeine synthesis in a coffee plant, incotine production in a tobacco plant, fruit ripening processes in an apple, tomato, pear, plum or peach plant, flower pigmentation in a rose, petunia, chrysanthemum or marigold plant or lignin production in a tobacco, aspen, poplar or pine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; c-raf; A-raf; B-raf; hammerhead ribozyme; hairpin ribozyme; traget; substrate; catalyst; modulation; expression; Rg gone; delivery; screening; identification; synthesis; deprotection; purification; cancer; inflammation; psoriasis; non-hepatic ascites; infection; genetic drift;
                                                                                                                                                                                                                                                                                                                                                            Skokut TA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ribozyme which modulates plant gene expression - preferably modulates expression of DELTA-9 desaturase or granule bound starch synthase in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
2.8%; Score 12; DB 1; Length 17;
Best Local Similarity 100.0%; Pred. No. 4.66+02;
Matches 12; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                      Mcswiggen JA, Merlo PAO,
Merlo DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 2 A; 5 C; 3 G; 0 T; 7 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 38; Page 86; 155pp; English.
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97US-0049002P.
97US-0051718P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV92424 standard; RNA; 17 BP
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                                                                                                                                                                                                                95US-0001135P.
                                                                                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC. (DOWC ) DOWELANCO.
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                                                                                                                                                                                                                                                                                                                                                            Zwick MG, Edington BE,
Young SA, Folkerts O,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       maize or canola.
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09-MAY-1997; 09-JUN-1997; 03-JUL-1997; JS-MAY-1998;

WO9850530-A2 Homo sapiens

AAV92424;

RESULT 922 AAV92424/c

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12-NOV-1998

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Gaps

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Human, hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Eming's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Alphel-Trenaunay-Weber syndrome; leukaemia; se; osteoporosis; DNAzyme; inozyme;

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids in samples, as amplification substrates, to nucleic acids in samples, as amplification substrates, to hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunospens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP-proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The concentration of skeletal muscle disorders. hGDMLP-1 may be used for diagnosing a disorders associated with the expression of hGDMLP-1, in particular heart of hGDMLP-1 is leaquence in the exemplification of the present sequence atta for this patent did not form part of the printed specification, but was obtained in electronic format directly from MIPO cat fitp.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                             New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                  Shannon ME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 12; DB 1; Length 17;
Pred. No. 4.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                  Chen W,
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100.0%; Pred. No. 3...
                                                                                                                                                                                                                                                                                                                                                                                  Rank DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; SEQ ID NO 1013; 214pp; English
                                                                                                                                                                                                                                                                                                                                                                                    Hanzel DK,
                                    2000US-0234687P.
2000US-0236359P.
2000US-0024263.
2001WO-US000661.
2001WO-US000663.
                                                                                                                                                                                                                                      30-JAN-2001; 2001WO-US000668
30-JAN-2001; 2001WO-US000668
30-JAN-2001; 2001WO-US000670
50-FEB-2001; 2001US-0266860P
                                                                                                                                                                                2001WO-US000665
                                                                                                                                                                                                                                                                                                                                                                                    Su Y, Ji Y, Penn SG,
                                                                                                                                                                                                                                                                                                                                             (AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-179446/23
                                                                                                                                     30-JAN-2001;
30-JAN-2001;
                                                                             04-OCT-2000;
30-JAN-2001;
                                                                                                                    30-JAN-2001;
                                                                                                                                                                                30-JAN-2001;
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Novel polynucleotide which down regulates expression of Ets-related gen useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.

Claim 4; Page 60; 149pp; English.

Jarvis T, Von Carlowitz I, Mcswiggen JA, Mclaughlin F, Randi AM;

16-MAY-2000; 2000US-00572021. 16-MAY-2001; 2001WO-US015866.

WO200188124-A2. Homo sapiens.

amberzyme.

22-NOV-2001

(RIBO-) RIBOZYME PHARM INC (GLAX) GLAXO GROUP LTD.

WPI; 2002-082995/11.

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The invention relates to a nucleic acid molecule (1) which down regulates coppession of an Ets-related gene (ERG). (I) is useful for treating capression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer. lymphoma, Ewing's sarcoma, melanoma, conditions selected from cancer. lymphoma, Ewing's sarcoma, melanoma, conditions aclerotis degeneration, arthritis, psoriasis, verruca vulgaris, angiofibroma, myopic degeneration, arthritis, psoriasis, verruca vulgaris, angiofibroma, of tuberous sclerosis, port-wine stains, Sturge weber syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies under conditions suitable for the treatment. Leukaemia or tumour conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ERG activity in a cell, by contacting (I) with RNA, in the presence of adivalent cation such as Mg2+. (I) is useful for diagnosis of conditions and cation such as Mg2+. (I) is useful for diagnosis of conditions and cation such as Mg2+. (I) is useful for diagnosis of conditions and cation such as hare homology with ERG gene or ERG fusion genes. Cargeting genes that share homology with ERG gene or ERG fusion genes. ABK7334-ABK22719 represent nucleic acids, including antisense and carged properserion cald molecules which regulate expression of ERG, and carged properserion acid molecules which regulate expression of ERG, and call call and call molecules which regulate expression of ERG, and call 
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.00.0%; Pred. No. 4.6e+02;
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Best Local Similarity 100.
Matches 12, Conservative
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13 AGAGGAGTCTCT RESULT 925

67

56 AGAGGAGTCTCT

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Gaps

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0; Indels

206 GAAAGCAGAGAA 217 Local Similarity 100.

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Query Match Matches 2 GAAAGCAGAGAA 13

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ABK18041/c ID ABK18041 standard; RNA; 17 BP.

Human ERG hammerhead ribozyme target sequence, Seq ID No 94.

RESULT 924
ABK17447/C
ID ABK17447 standard; RNA; 17 BP.
XX
AC ABK17447;
XX
DT 09-APR-2002 (first entry)
XX
DE Human ERG hammerhead ribozyme ta

ABK18041;

(first entry) 09-APR-2002 Human ERG hammerhead ribozyme target sequence, Seq ID No 688.

Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; center; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour andiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss; Osler-Weber-rendu syndrome; leukaemia; osteoporosis; DNAzyme; inozyme;

Homo sapiens.

WO200188124-A2.

22-NOV-2001.

16-MAY-2001; 2001WO-US015866.

L6-MAY-2000; 2000US-00572021.

RIBO-) RIBOZYME PHARM INC GLAX) GLAXO GROUP LTD Jarvis I, Von Carlowitz I, Mcswiggen JA, Mclaughlin F, Randi AM;

WPI; 2002-082995/11.

Novel polynuclectide which down regulates expression of Ets-related genuseful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.

Claim 4; Page 71; 149pp; English.

The invention relates to a nucleic acid molecule (I) which down regulates corpression of an Ets-related gene (ERG). (I) is useful for treating carcoms, machines selected from cancer, lymphoma, Ewing's sarcoms, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, tumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiofibroma of tubberous sclerosis, port-wine stains, Sturge weber syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies to the treatment. The method comprises the use of one or more therapies conditions suitable for the treatment. Leukaemia or tumour conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ERG activity in a cell, by contacting (I) with RNA, in the presence of advalent cation such as Mg2+. (I) is useful for diagnostic coll to cation such as Mg2+. (I) is useful for diagnostic coll to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically cargeting genes that share homology within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically cargeting genes that share homology within diseased cells or conditions and cargeting genes that share homology within diseased cells or condition or enzymatic nucleic acid molecules which regulate expression of ERG, and cargeting decent cargeting massed acids, including antisense and cargeted profile acid molecules which regulate expression of ERG, and conditions and called PCR primers of the invention

Sequence 17 BP; 4 A; 6 C; 3 G; 0 T; 4 U; 0 Other;

Gaps ö Length 17; Indels ö 2.8%; Score 12; DB 1; Le 100.0%; Pred. No. 4.6e+02; ive 0; Mismatches 0; 2.85, 100.0%; Fre Conservative Best Local Similarity Matches 12; Conserv Query Match

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14 AGAGGAGTCTCT

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RESULT 926

BP. ABK18042 standard; RNA; 17 **ABK18042**

ABK18042;

(first entry) 09-APR-2002 Human ERG hammerhead ribozyme target sequence, Seq ID No 689.

Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; sturge Weber syndrome; kippel-Trenaunay-Weber syndrome; leukaemia; se; Osler-Weber-rendu syndrome; leukaemia; osteoporosis; DNAzyme; inozyme; amberzyme.

Homo sapiens.

WO200188124-A2.

22-NOV-2001

16-MAY-2001; 2001WO-US015866.

16-MAY-2000; 2000US-00572021.

(RIBO-) RIBOZYME PHARM INC. (GLAX) GLAXO GROUP LTD.

Randi AM; Mclaughlin F, Mcswiggen JA, Von Carlowitz I, WPI; 2002-082995/11. Jarvis T,

Novel polynucleotide which down regulates expression of Ets-related gen useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.

Claim 4; Page 71; 149pp; English.

The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiogenesis, diabetic retinopathy, macular degeneration, cumour supporting angiofibroma of tuberous sclerosis, port-wine stains, Sturge weber syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more theraples to under conditions suitable for the treatment. Leukaemia or tumour conjunction with one or more of other theraples such as radiation or conjunction with one or more of other theraples such as radiation or conjunction with one or more of other theraples such as radiation or cell, by contacting (I) is useful for reducing ERG activity in a cell of seases related to the expression of ERG, and as diagnostic tool to diseases related to the expression of ERG, and as diagnostic coll to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically crafting genes that share homology within ERG gene or ERG fusion genes. ABK13354-ABK22719 represent nucleic acids, including antisense and carged molecules which regulate expression of ERG, and crafted PCR primers of the invention

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Gaps

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The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, melanoma, myopic degeneration, arthritis, psoriation, vernuca vulgaris, angiofibroma of tuberous sclerosis, portwine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-rendu wend reating a patient having a condition associated with the level of ERG, by contacting calls of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more theraples under conditions suitable for the treatment. Leukaemia or tumour conditions suitable for the treatment. Leukaemia or tumour angiogenesis is treated by administering (I) to the patient in conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ERG activity in a coll, by contacting the cell with RNA, in the presence of a divalent cation such as Mg2+. (I) lis useful for diagnosis of conditions and diseases related to the expression of ERG, and as diagnostic tool to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human, hammerhead ribozyme, cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerzy; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous solerosis; port-wine stain; wound healing; Sturge Weber syndrome; kippel-Trenaunay-Weber syndrome; leukaemia; ss; Osler-Weber-rendu syndrome; leukaemia; osteoporosis; DNAzyme; inozyme;
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                                                                    Length 17;
                                                                                                                                            0; Indels
Sequence 17 BP; 4 A; 7 C; 2 G; 0 T; 4 U; 0 Other;
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                                                                    Score 12; DB 1; L. Pred. No. 4.6e+02;
                                                                                                                                        Mismatches
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                                         2.8%; Scc.
100.0%; Pre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABK18967 standard; RNA; 17 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (RIBO-) RIBOZYME PHARM INC
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                                                                                                                                        Conservative
                                                                                                                                                                                                             56 AGAGGAGTCTCT 67
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                                                                                              Local Similarity
es 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABK18967;
                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 927
                                                                                                                                            Matches
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XXX ABK1
XXX ABK1
XXX ABK1
XXX ABK1
XXX ABK1
XXX BUIMR
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The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Bwing's sarcona, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, neovascular glaucoma, myopic degeneration, arthritis, psoriasis, verruca vulgaris, angiofibroma of tuberous sclerosis, port-wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-rendu syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by conteacting cells of the patient with (I) under conditions suitable for the treatment. Demethod comprises the use of one or more therapies under conditions suitable for the treatment. Leukaemia or tumour
                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel polynucleotide which down regulates expression of Ets-related genuseful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human, hammerhead ribozyme, cytostatic, antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide, osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous selerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; se; osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme;
                 the presence of ERG RNA in a cell. (I) is useful for specifically targeting genes that share homology with ERG gene or ERG fusion genes. ABK17354-ABK22719 represent nucleic acids, including antisense and enzymatic nucleic acid molecules which regulate expression of ERG, and related PCR primers of the invention
examine genetic drift and mutations within diseased cells or to detect
                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                      2.8%; Score 12; DB 1; Length 17; 100.0%; Pred. No. 4.6e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human ERG Amberzyme target sequence Seg ID No 1872.
                                                                                                                                             Sequence 17 BP; 4 A; 5 C; 3 G; 0 T; 5 U; 0 Other;
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ID ABK19225 standard; RNA; 17 BP.
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                                                                                                                                                                      Query Match
Best Local Similarity 100..
Best Local 12; Conservative
                                                                                                                                                                                                                                                                           67
                                                                                                                                                                                                                                                                           56 AGAGGAGTCTCT
                                                                                                                                                                                                                                                                                                                    16 AGAGGAGTCTCT
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anglogenesis is treated by administering (I) to the patient in conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ERG activity in a cell, by contacting the cell with (I). (I) is useful for cleaving RNA of ERG gene, by contacting (I) with RNA, in the presence of a divalent cation such as Mg4. (I) is useful for diagnosis of conditions and diseases related to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically targeting genes that share homology with ERG gene or ERG fusion genes. ABKI7354-ABK22719 represent nucleic acids, including antisense and enzymatic nucleic acid molecules which regulate expression of ERG, and related PCR primers of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (S1, ABB83999), a sequence having 65% sequence identity to (S1),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human, POSH1 1, SH3 domain, POSH-like signalling protein 1, oncogene, Rho GTPase; signal transduction, gene expression, cancer, vaccine;
                                                                                                                                                                                                                                                                                                                        0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3xample 2; SEQ ID NO 1753; 60pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                               2.8%; Score 12; DB 1; Length 17; 100.0%; Pred. No. 4.6e+02; trive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human POSHL1 scanning oligonucleotide SEQ ID NO 1753.
                                                                                                                                                                                                                                                   Sequence 17 BP; 4 A; 6 C; 3 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-JAN-2001, 2001WO-US000663.
30-JAN-2001, 2001WO-US000665.
30-JAN-2001, 2001WO-US000665.
30-JAN-2001, 2001WO-US000667.
30-JAN-2001, 2001WO-US000667.
30-JAN-2001, 2001WO-US000668.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene therapy; transgenic; ss.
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                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                            56 AGAGGAGTCTCT 67
                                                                                                                                                                                                                                                                                                                                                                                               17 AGAGGAGTCTCT 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-684061/74.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shannon M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABV91040;
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ABV91040/c
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fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as adometreem components of the signal transduction pathway. (I) is useful ac downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) encoding (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHL1 including disease and treating caused by altered expression of human POSHL1 including disease and (II) is treating cancer, they useful in the development of vaccines and (II) is cusful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonoclocited useful in examples of the invention. Note: The present sequence did not form part of the print by the Buropean Patent Office
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human, pharmacological; hypotensive; antilipaemic; vasotropic; laxative; dermatological; antidepressant; tranquilliser; antiinflammatory; eczema; antiluler; antimigrafine; neuroprotective; antiparkinsonian; analgesic; yanaecological; virucide; vulnerzry; antiarthitic; antipsoriatic; col; yanaecological; virucide; vulnerzry; antiarthitic; antipsoriatic; col; antimicrobial; cytostatic; litholytic; pathological disorder; depression; antimicrobial; cytostatic; litholytic; pathological disorder; depression; whorefulle dysfunction; anxiety; stress; inflammatory bowel syndrome; we rectile dysfunction; anxiety; stress; inflammatory bowel syndrome; we constibation; headeache; seizure; multiple sclerosis; polymyositis; fibromyalgia; Parkinson's disease; anyotrophic lateral solerosis; trauma; chronic pain; pre-menstrual syndrome; sinusitis; carpal tunnel syndrome; chronic fatigue syndrome; rosacea; arthritis; psoriasis; prostatis; millammation; heat burn; infection; colon cancer; malignant melanoma; we skin disorder; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Composition with a modified oligonucleotide useful for treating a patient with a pathological disorder such as abnormal appetite, hypertension, eczema, anxiety, stress, and cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 2 A; 6 C; 6 G; 3 T; 0 U; 0 Other;
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Claim 17; Page 9; 173pp; English.

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The present invention describes a composition (I) suitable for administration in a mammal, which comprises a modified oligonucleotide dainistration in a mammal, which comprises a modified oligonucleotide scontaining 7 or more contiguous ribose groups linked by achiral 5'-3' internucleoside phosphate linkages, where the modified oligonucleotide is complementary to a region of a gene comprising (II) and (2) a cosmetic composition comprising (II), where the modified oligonucleotide is complementary to a region of a gene associated with a skin disorder. (I) and (II) and ill can have hypotensive, antilinflammatory, antilucer, laxative, antimigration, corrective, antilinflammatory, antilucer, laxative, antilinflammatory, antilucer, laxative, antimigration, corrective, antilarthritic, antilosorial, symaeological, virucide, ranguilliser, antilarthritic, antipsoriatic, antimicrobial, cytostatic and litholytic activities. (I) can be used for treating a patient with a pathological disorder selected from abnormal appetite, hypertension, corrective, antistriptical and antipsorial companies, inflammatory bowel syndrome, ulcerative colitis, Crohn's disease, renal stones, gall stones, constipation, colds, migrain, per-menstrual syndrome, trauma, carpal tunnel syndrome, colitis, chronic pain, per-menstrual syndrome, trauma, carpal tunnel syndrome, colitis, chronic fatigue syndrome, rosacea, arrhitis, poriasis, prosaceis, sinalignant composition is desease, anyotrophic lateral solerome, malignant and malignant nasal polyps. The nutritional supplement is composition is useful for improving the appearance of the skin in an individual with a skin disorder. Acressor of the present invention composition in the exemplification of the present invention
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0; Gaps Query Match
2.8%; Score 12; DB 1; Length 17;
Best Local Similarity 100.0%; Pred. No. 4.6e+02;
Matches 12; Conservative 0; Mismatches 0; Indels Sequence 17 BP; 2 A; 4 C; 9 G; 2 T; 0 U; 0 Other;

146 GGTGGAGGCCGG 157 4 GGTGGAGGCCGG 15 ઠે 쉱

RESULT 931

ABT39673 standard; DNA; 17 BP. 12-JUN-2003 (first entry) ABT39673;

Tumour suppression related human fukutin oligo SEQ ID No 5310.

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.

Homo sapiens.

WO2003025175-A2.

27-MAR-2003

17-SEP-2001; 2001FR-00011978.

17-SEP-2002; 2002WO-IB004208

(MOLE-) MOLECULAR ENGINES LAB

Tuijnder M; relerman A, Amson R,

WPI; 2003-313353/30

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.

Disclosure; Page 654; 720pp; French.

The invention relates to a novel isolated 17 mer nucleic acid sequence, c given in the specification, a sequence containing at least 15 consecutive c given in the specification, a sequence with, after optimal ancelorated sequence with, after optimal alights to the 17 mer sequence, a sequence that alignment, at least 80 % identity to the 17 mer sequence that the alights to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated mucleic cide of the invention are useful as probes and primers for detecting, caids of the invention are useful as probes and primers for detecting, component of a gene chip, in vitro as (anti) sense reagents, and for production of recombinant polypeptides. Any of the mucleic acids, component of a gene chip, in vitro as (anti) sense reagents, and for production of recombinant polypeptides. Any of the mucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for vector or antibodies directed against the polypeptides are useful for components of parameterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and degeneration, specifically cancer but also Alzheimer's disease and diseases. The polypeptides can also be used to generate antibodies, and colings. The nucleic acid sequences of the invention can be used in gene therapy. This polymucleotide sequence represents a tumour suppression contexts.

Sequence 17 BP; 6 A; 7 C; 2 G; 2 T; 0 U; 0 Other;

Gaps .; o Query Match
2.8%; Score 12; DB 1; Length 17;
Best Local Similarity 100.0%; Pred. No. 4.66+02;
Matches 12; Conservative 0; Mismatches 0; Indels

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ACA07722 standard; RNA; 17 BP. RESULT 932 ACA07722/c

03-JUN-2003 (first entry) ACA07722;

NFKB sub-unit modulating zinzyme substrate #121.

G-cleaver, amberryme; cancer; REL-A activity; breast cancer; human; lung cancer; amberryme; cancer; REL-A activity; breast cancer; human; lung cancer; amberryme; colorectal cancer; brain cancer; human; cesophageal cancer; colorectal cancer; brain cancer; prostate cancer; cervical cancer; head and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; disablatin; methotraxate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoin arthritis; restenosis; Crohn's disease; obesity; lachaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/Graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammatory bowel disease; infection; ss. snzymatic nucleic acid, nuclear factor kappa B; NFKB; inozyme; zinzyme;

Homo sapiens

JS2002177568-A1.

28-NOV-2002

23-MAY-2001; 2001US-00864785

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The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFKB), where (I) is an inozyme, G-leaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG^2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glicma or multidrug resistant cancer; The method involves use of other drug chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, detrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, detrexate, cyclophosphamide, doxorubin, gluorouracil carboplatin, detrevate, cyclophosphamide, doxorubin, gluorouracil carboplatin, detrevate, cyclophosphamide, doxorubin, gluorouracil carboplatic, and antiense nucleic central nervous system (CNS) and mycoardial, glumentol antiense or antient cyclophosphamide, doxorubin, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic ciral molecule
                                                                                                                                                                                                                                                                                                                       Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.
                                                                                                                                                                                                                               Stinchcomb DT, Mcswiggen J, Draper KG
                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 3; Page 39; 72pp; English.
  92US-00987132.
94US-00245466.
94US-00291932.
96US-00777916.
                                                                                                                           STINCHCOMB D T.
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                                                                                                                                                                                                                                                                            WPI; 2003-340953/32.
                                                                                                                                                                            DRAP/) DRAPER K G.
                                                                                                                                                      MCSWIGGEN
07-DEC-1992;
18-MAY-1994;
15-AUG-1994;
23-DEC-1996;
                                                                                                                              (STIN/)
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2.8%; Score 12; DB 1; Length 17; 100.0%; Pred. No. 4.6e+02; ve 0; Mismatches 0; Indels Sequence 17 BP; 2 A; 11 C; 3 G; 0 T; 1 U; 0 Other; 2.5°, 100.0%; Pre-Local Similarity 100 nes 12; Conservative Query Match Best Loc Matches

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0; Gaps

ACA07649 standard; RNA; 17 BP RESULT 933

03-JUN-2003 (first entry) ACA07649;

NFKB sub-unit modulating zinzyme substrate #48.

Enzymatic mucleic acid; muclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; oesophageal cancer; stomach cancer; bladder cancer; pancreatic cancer; head and neck cancer; ovarian cancer; melanoma; llymphona; glinoma; multidrug resistant cancer; melanoma; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate;

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The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor regulates expression of a sequence encoding a subunit of nuclear factor configuration. The enzymatic nucleic acid molecule is adapted to treat configuration. The enzymatic nucleic acid molecule is adapted to treat configuration. The enzymatic nucleic acid molecule is adapted to treat configuration. The enzymatic nucleic acid molecule are ussociated with the level of RBL-A. CC treating a patient having a noomitising a sequence of RBL-A gene, in the presence of a divalent cation, especially Mg^2+. The enzymatic and the presence of a divalent cation, especially Mg^2+. The enzymatic and contisted molecules are useful for treating breast lumg, contisted, brain, necknows, lymphoma, glioma or multidural resistant cancer. The method involves use of other drug cervical, brain and national antibodies, RBL-A-specific inhibitors or therapies such as monoclonal antibodies, RBL-A-specific inhibitors or chemotherzapy including paclitaxel, doctaxel, cisplain, methotizexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, condimined are also useful for treating inflammatory disease such as colored acid activities, restenosis, asthma, Crohm's disease, diabetes, obesity, autoimmune disease, luus, multiple sclerosis, transplant/graft crifection, gene therapy applications, miltiple sclerosis, transplant/graft crifection, proper energy applications, inflammatory bowel disease or confident and myocardial), glomerallonephritis, and myocardial), glomerallonephritis, and myocardial miltiple sclerosis a novel enzymatic confidence represents the substrate of a novel enzymatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.
                    rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy, autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonophritis; allergic airway inflammation; inflammatory bowel disease; infection; ss.
radiation therapy; inflammatory disease; asthma; diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 4 A; 4 C; 6 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Draper KG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 3; Page 38; 72pp; English
                                                                                                                                                                                                                                                                                                                                        92US-00987132.
94US-00245466.
94US-00291932.
96US-00777916.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Stinchcomb DT, Mcswiggen J,
                                                                                                                                                                                                                                                                                            23-MAY-2001; 2001US-00864785
                                                                                                                                                                                                                                                                                                                                                                                                                                                                (STIN/) STINCHCOMB D T.
(MCSW/) MCSWIGGEN J.
(DRAP/) DRAPER K G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      infection. This sequen
nucleic acid molecule
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18-MAY-1994;
15-AUG-1994;
23-DEC-1996;
                                                                                                                                                   Homo sapiens.
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Score 12; DB 1; Length 17; Pred. No. 4.6e+02; 1; Mismatches 0; Indels ВЪ 2.8%; ACC64123/c ID ACC64123 standard; DNA; 17 XX 268 ACCTGGAGCAGG 279 Query Match Best Local Similarity 91.7 Matches 11, Conservative 2 AccudeAdcade 13 8

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Gaps ÷ Distinguishing multiple alleles and identifying new alleles - by single-strand conformation polymorphism technique using specific gel electrophoresis conditions.

Disclosure; Page 19; 36pp; English.

White MB

Carrington M,

Mann D,

WPI; 1993-017809/02. Dean M,

(USSH) US DEPT HEALTH & HUMAN SERVICE.

91US-00751892 91US-00751892

29-AUG-1991; 29-AUG-1991;

01-DEC-1992.

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The present invention relates to murine oligonuclectides (ACC62754-ACC68806), which are associated with tumour suppression, tumour reversion, apoptosis and virus resistance. The oligonucleotides are useful as (1) as probes and primers for detecting, identifying, quantifying and/or amplifying nucleic acid, e.g. as one component of a gene chip; in vitro as (anti)sense reagents; and (2) for production of recombinant polypeptides. The oligonucleotides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies
                                                                                                     Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease;
                                                                   Aurine oligonucleotide associated with tumour supression, SEQ ID 1370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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2.8%; Score 12; DB 1; Length 17;
Best Local Similarity 100.0%; Pred. No. 4.6e+02;
Matches 12; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 6 A; 1 C; 7 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Tuijnder M;
                                                                                                                                                                                                                                                                                                                                                                                             (MOLE-) MOLECULAR ENGINES LAB
                                                                                                                                                                                                                                                                                                                     7-SEP-2002; 2002WO-IB004210.
                                                                                                                                                                                                                                                                                                                                                         17-SEP-2001; 2001FR-00011979
                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    vPI; 2003-333167/31.
                                                                                                                                                                     schizophrenia; ss
                                                                                                                                                                                                                                           102003025176-A2
                                                                                                                                                                                                         fus musculus
                                     01-JUL-2003
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The oligomer AGI represents a probe for DQAI alleles 0101, 0102 and 0103

and is used to distinguish multiple alleles of a gene of the immunoglobin

supergene family. The DNA encoding the gene of interest in a sample is

amplified and then denatured. The amplified DNA is then separated on a

mon-denaturing polyarylamide gel consisting of 5 percent bis-acrylamide

condenaturing polyarylamide gel consisting of 5 percent bis-acrylamide

with 0-10 percent glycerol, and the presence or absence of DNA bands

consisting hybridisation is detected. Before amplification of the gene, the

alleles may be divided into subsets by oligomucleotide hybridisation.

consisting single stranded conformation polymorphism (SSCP) multiple alleles

con new alleles identified. The method may be used in studying genetic

and new alleles identified. The method may be used in studying genetic

consisting which correlate with the presence of disorders such as cystic

contransplantation. The SSCP method has been used for detection of mutant

contransplantation. The SSCP method has been used for detection of mutant

contransplantation. The SSCP method has been used for detection of mutant

contransplantation in the presence of disorders such as cystic

contransplantations to prevent clashes with ongoing US granted patent

contransplantations to prevent clashes with ongoing US granted patent

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intracellular cyclic nucleotide level modulation; cAMP; cGMP; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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Amplification; conformation polymorphism; SSCP; DQ-alpha; DQ-beta; cystic fibrosis; neurofibromatosis; ss.

USN7751892-N

Synthetic.

DQA1 probe AG1, for alleles 0101, 0102 and 0103.

(revised)
(first entry)

17-DEC-2001 12-MAY-1993

AAQ34452;

AAQ34452 standard; DNA; 18 BP.

RESULT 935 AAQ34452

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RESULT 938
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                                                                                                                                                                                                                                                 This sequence is a PCR primer for DNA encoding the human phosphodiesterase 8 (PDE8) of the invention. The phosphodiesterase genes and polypeptides are used to develbo products for treating conditions in which cyclic nucleotide pathways are aberrant and for modulation of intracellular cyclic nucleotide levels. The PDE8 polypeptides exhibit high affinity for hydrolysis of both cAMP and cGMP but relatively low sensitivity to enzyme inhibitors specific for other PDE families. The PDE8A polypeptides and polymucleotides can be used for identifying their specific binding partners. The products can provide approaches for treating conditions in which cyclic nucleotide pathways are aberrant as well as conditions in which modulation of intracellular cAMP and/or cGMP levels in certain cell types is desirable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BLM gene cluster; bleomycin gene cluster; polyketide metabolite;
bleomycin; bleomycin analogue; holo-carrier protein; thiazolidine;
thiazoline; bithiazoline; microbial metabolite; sugar; PCR primer; ss.
                                                                                                                                               New isolated phosphodiesterase genes and polypeptides for identifying specific binding partners.
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100.0%; Pred. No. 5.1e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 18 BP; 4 A; 7 C; 2 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                   Example 3; Page 14; 80pp; English.
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AAA58509 standard; DNA; 18 BP.
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05-FEB-1999; 99US-0118848P.
05-JAN-2000; 2000US-00477962.
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                                                                                                          WPI; 1999-277645/23
                    (ICOS-) ICOS CORP
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                                                                Loughney K;
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Best Local Si
Matches 12,
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SX T T T T X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B
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The sequence is a probe used in a Tagman real-time PCR experiment used to demonstrate that mice deficient of type I cytokine receptor, mTCCR, are impaired in their ability to mount a Th1 response. The invention relates to methods of modulating the differentiation of T-cells into the Th2 subtype instead of the Th1 subtype, by administering a modulator of TCCR (e.g. an antagonist) to enhance, stimulate or potentiate T-cell differentiation, or using TCCR polypeptide or its agonists to prevent, inhibit or attenuate T-cell differentiation. Th1 mediated disease in mammal can be treated by administering a TCCR antagonist and Th2 diseases by administering a TCCR antagonist and Th2 diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                    (ORR9) B to 41 of the BLM (Bleenwyin) gene cluster. The proteins encoded by the gene cluster are useful for producing peptides and/or polyketide metabolites, especially bloomycin or bleomycin analogues. They are also useful for chemically modifying biological molecules to produce branched methyl groups, and for coupling amino acids and fatty acids. They may be reacted with an apo-carrier protein and coenzyme A to produce a holocarrier protein. The BLM gene cluster or catalytic domains can be used bithiazoline and bithiazoline, containing microbial metabolites. The BLM gene cluster way also be used to produce sugars
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mouse; type-I cytokine receptor; TCCR; T-cell differentiation; Th1; Th2; agonist; antagonist; autoimmune inflammatory disease; allograft rejection; multiple sclerosis; inflammatory bowel disease; insulin-dependent diabates mellitus; infectious disease; human immunodeficiency virus; allergic disorder; asthma; allergic thinitis; HIV; probe; mRPL19; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Modulating T-cell differentiation and cytokine release profiles into Th1 and Th2 subtypes, for treating immune-related diseases in mammals, by administering modulator of type I cytokine receptor (TCCR).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
primers AAA58474-A58541 were used to amplify open reading frames
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 18; 5.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 18 BP; 1 A; 8 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 2.8%; Score 12; DB Best Local Similarity 100.0%; Pred. No. 5.1 Matches 12; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 12; Fig 19; 126pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAS03269 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-OCT-2000; 2000WO-US028827.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-0160542P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mouse mRPL19 Taqman probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    326 GGCGGCGGACGA 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GGCGCGGACGA 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-308474/32.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus
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AAS07305 standard; DNA; 18 BP

RESULT 940 AAS073

3 GTCTCTGCACTA 14

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(first entry)

12-SEP-2001

AAS07305;

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The present invention relates to human phosphodiesterase 8 (PDE8) (AAC63695 and AAD28256). Phosphodiesterases hydrolyse 3', 5' cyclic uncleotides to their respective mucleoside 5' monophosphates. PDE8 may be used in the systematic analysis of the structure and function of PDE8, and for the identification of molecules with which PDE8 will interact. PDE8 coding sequence may be used in hybridisation assays to detect the capacity of cells to express PDE8, and as a basis for diagnostic methods useful for identifying a genetic alteration in a PDE8 locus that underlies a disease state or states. The human PDE8 gene has been localised to chromosome 6p36-27. The present sequence is a PCR primer used to isolate the coding sequence of human PDE8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; PDE8; phosphodiesterase 8; chromosome 6p26-27; PCR primer; ss.
rejection and autoimmune inflammatory diseases, such as allergic encephalomyelitis, multiple sclerosis, insulin-dependent diabetes mellitus, autoimmune vveoretinitis, inflammatory bowel disease or autoimmune thyroid diseases. The mediated diseases include infectious diseases, such as Leishmania major, Mycobacterium leprae, Candida albicans, Toxoplasma gondii, respiratory syncytial virus and human immunodeficiency virus (HIV) and allergic disorders, such as asthma, allergic rhinitis, dermatitis and vernal conjunctivitis
                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New phosphodiesterase BA (PDEBA) polypeptides useful used in the systematic analysis of the structure and function of PDEB, and for identifying molecules with which PDEBA will interact.
                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                              Query Match 2.8%; Score 12; DB 1; Length 18; Best Local Similarity 100.0%; Pred. No. 5.1e+02; Matches 12; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 18 BP; 4 A; 7 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                           Sequence 18 BP; 0 A; 7 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; Col 10; 37pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                    AAC66689 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98US-00174437.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human PDE8 PCR primer W48A9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                246 TICCCGGGCTCG 257
                                                                                                                                                                                                                                                                                                                                   1 Trecesseries 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ICOS-) ICOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US6133007-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAC66689;
                                                                                                                                                                                                                                                                                                                                                                                             RESULT 939
AAC66689
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The sequence represents a sequencing primer used to sequence a genomic clone from Cochliobolus heterostrophus which contains the CPS1 and TES1 peptide syntherase genes. CPS1 is an enzyme thought to be involved in the production of peptide toxins, which are involved in the pathogenic infection of corn crops. The nucleic acids and proteins can be used as tragers for anti-fungal compounds to prevent fungal corn infection and the nucleic acids can be used in gene therapy to alter the biosynthetic pathway for the peptide toxins to lower the pathogenicity of the fungi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                   New isolated nucleic acid molecule from a plant pathogen useful in
preventing plant pathogenic infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
2.8%; Score 12; DB 1; Length 18;
Best Local Similarity 100.0%; Pred. No. 5.1e+02;
Matches 12; Conservative 0; Mismatches 0; Indels
                                                                                                 CPS1; peptide synthetase; peptide toxin; fungal pathogen; corn crop infection; 88; sequencing primer; FP8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human chromosome 1p36-35 PCR primer SEQ ID NO:1779.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 18 BP; 4 A; 4 C; 5 G; 5 T; 0 U; 0 Other;
                                                                    CPS1/TES1 genomic DNA sequencing primer FP8
                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 54; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABL44735 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                 (CORR ) CORNELL RES FOUND INC.
                                                                                                                                                                                                                                                                                                                              Lu S;
                                                                                                                                                                                                                                        22-NOV-2000; 2000WO-US032227.
                                                                                                                                                                                                                                                                    99US-00448215
                                                                                                                                                Cochliobolus heterostrophus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           261 ACGGTGCACCTG 272
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                                                                                                                                                                                                                                                                                                                              Yoder OC, Turgeon BC,
                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-367672/38.
                                                                                                                                                                            WO200138489-A2
                                                                                                                                                                                                                                                                    23-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-APR-2002
                                                                                                                                                                                                           31-MAY-2001
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ABL44735/c
CXXXEXEXEXXXXXXX
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Gaps

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2.8%; Score 12; DB 1; Length 18; 100.0%; Pred. No. 5.1e+02; rative 0; Mismatches 0; Indels

Local Similarity 100.

Matches

Query Match

62 GTCTCTGCACTA 73

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The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are maxed in each of the multiwell plates; (b) a primer designed based on the chromosome marker captened is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant amplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the maximum in the specified discrimination Nos. to array the multiwell constrainination Nos. to array the multiwell the maximation Nos. to make discrimination Nos. to array the multiwell cand lateral directions; (f) the mixed clones are cultured and the capultant cultures are mixed respectively in each wells of longitudinal cand lateral directions; (f) the mixed clones are cultured and the resultant cultures are amplified by using the above primer; (g) signals care detected from the amplified products; (h) the clones in the multiwell plates are specified from the amplified products; (h) the clones in the multiwell constituted as the positions on the chromosome and arrayed. The constituted as the positions on the chromosome and arrayed. The constituted mixed for muman chromosome 21022.1, which are specified for use in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 1 A; 4 C; 7 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                    Claim 4; Page 39; 528pp; Japanese.
                                                                                                                                    10-MAR-2000; 2000JP-00066716.
                                                                                          12-MAR-2001; 2001JP-00068285
                                                                                                                                                                                    (RIKA ) RIKAGAKU KENKYUSHO (GENO-) GENOTEX YG.
                                                                                                                                                                                                                                                                                                       Arraying genome clones.
                                                                                                                                                                                                                                                       WPI; 2002-144136/19.
JP2001321190-A.
                                             20-NOV-2001
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0; Gaps
Query Match 2.8%; Score 12; DB 1; Length 18; Best Local Similarity 100.0%; Pred. No. 5.1e+02; Matches 12; Conservative 0; Mismatches 0; Indels
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ABS68429 standard; DNA; 18 BP. RESULT 942 ABS68429

19-NOV-2002 (first entry) ABS68429;

Sequencing primer #20 for fungal DNA flanking REMI insertion site.

Fungal pathogen; peptide synthetase gene cluster; iron reductase; permease; major facilitator superfamily transporter; MFS transporter; anti-fungal agent; fungicide; pathogenic fungi; plant pathogen; CFS1; animal pathogen; fungal infection; wild grass; cereal; corn; mycocide; leaf spot maize; immunocompromised vertebrate; pneumonia; arthritis; miliary disease; bone infection; joint infection; skin disease; aesophagitis; vaginitis; onychomycosis; inflammation; urinary tract; kidney; liver; brain; gastrointestinal tract; lung; fungicidal; mycocidal; antiarthritic; antinflammatory; dermatological; COA ligase; sequencing; primer; ss.

Cochliobolus heterostrophus Synthetic

21-NOV-2001; 2001WO-US043381. WO200242444-A2 30-MAY-2002.

22-NOV-2000; 2000US-0252649P. 22-NOV-2000; 2000US-0252732P.

(SYGN) SYNGENTA PARTICIPATIONS AG. (CORR) CORNELL RES FOUND INC. (VDDE/) YODER O. (TURG/) TURGEON B G. (LUSS/) LU S.

L S; Turgeon BG, foder 0,

NPI; 2002-666824/71.

Nucleic acid molecules comprising fungal, e.g. Cochliobolus heterostrophus, genes from a peptide synthetase gene cluster, useful for identifying anti-fungal agents for treating fungal infections such as pneumonia and arthritis.

Example 1; Page 188; 315pp; English.

The present invention relates to nucleic acid molecules comprising
fungal, e.g. Cochliobolus heterostrophus, genes from a peptide synthetase
concluster, encoding e.g. an iron reductase and/or a permesse, or a
major facilitator superfamily (MFS) transporter protein. The
polynucleotides and polypeptides are useful for identifying a novel
thinbitors of gene products that are useful as fungicides or
covel inhibitors of gene products that are useful as fungicides or
mycocides. Anti-fungal agents identified using the polynucleotide and
covel inhibitors of gene prowth of pathogenic fungi. The fungal
consideration of the invention, and antisense DNA are useful as
fungicides to suppress the growth of pathogenic fungi. The fungal
consideration include plant pathogens such as Septoria trici, or Cochliobolus
considerates, adisease called leaf spot maize caused by the pathogen C
treat a disease called leaf spot maize caused by the pathogen C
treat a disease called leaf spot maize caused by the pathogen C
treating fungal infections of vertebrates, including immunocompromised
construction from skin disease, asophagitis, vaginitis, onychomycosis,
and inflammation of the urinary tract, kidney, liver, brain,
construction site in the examples of the present invention
construction
construction site in the examples of the present invention

Sequence 18 BP; 4 A; 4 C; 5 G; 5 T; 0 U; 0 Other;

Gaps ő Query Match
2.8%; Score 12; DB 1; Length 18;
Best Local Similarity 100.0%; Pred. No. 5.1e+02;
Matches 12; Conservative 0; Mismatches 0; Indels

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ABA03691 standard; DNA; 18 BP. ABA03691; ABA03691
ID ABA0
XX
XX
AZ ABA0
XX
DT 18-F
XX

RESULT 943

18-FEB-2002 (first entry)

HSV-tk gene-del PCR primer TrTkl.

Cytostatic; antitumour; gene therapy; thymidine kinase; tk;

Triticum aestivum

WO200299111-A2

12-DEC-2002

07-JUN-2001; 2001US-0296159P. 07-JUN-2002; 2002WO-CA000834

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The invention relates to a polynucleotide encoding a thymidine kinase (tk), where the tk coding region does not contain a functional splice acceptor and/or splice donor site. The polynucleotide and the protein that it encodes are useful for destroying cells. The polynucleotide is introducing into the cells, allowing the cells to express tk. The cells are then contacted with a substantially non-toxic agent which is converted by tk into a toxic agent. The non-toxic agent which is converted by tk into a toxic agent. The non-toxic agent wish as converted by tk into a toxic agent. The non-toxic agent wish of the cells that are lost of uranosyl) -2-iddouradil, ara-A, ara | 1.-beta-D arabino furanosyl -2-iddouradil, ara-A, ara | 1.-beta-D arabino furanosyl thymine, 5-iddouradil, ara con bromovinyl deoxyuridine, idoxuridine, AZT, AIV, dideoxycytidine, Ara C or bromovinyl deoxyuridine therpy, and for manufacturing a sedicament for destroying cells in a patient. The polymucleotide is used to destroy cells that are, or have the potential to become, cancer cells. The polymucleotide does not contain a splice donor and/or splice acceptor site, and so there is no undesirable splicing, which would lead to the production of an aberrant form of the thymidine kinase gene. Thus a greater proportion of transduced target cells correctly express tk. The present sequence is a primer used to selectively amplify the deleted form of the herpes simplex virus (HSV)-tk gene using a 5' primer, which spans the truncation point
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel polynucleotide comprising a thymidine kinase coding region encoding thymidine kinase, which does not contain a functional acceptor and/or splice donor site, useful for gene therapy techniques.
splice acceptor site; splice donor site; cell destruction; cytostatic; cancer; herpes simplex virus; HSV; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                        (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sxample 3; Page 59; 103pp; English.
                                                                                                                                                                                                                                                                  .3-APR-2001; 2001WO-GB001640.
                                                                                                                                                                                                                                                                                                                     .3-APR-2000; 2000GB-00008966.
                                                                                 Herpes simplex virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-026030/03
                                                                                                                                                             40200179502-A2
                                                                                                                                                                                                                                                                                                                                                                                                                          Apperley JF,
                                                                                                                                                                                                                25-OCT-2001
                                                                                                             Synthetic
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                                                             Gaps
                         Score 12; DB 1; Length 18; Pred. No. 5.1e+02; 0; Mismatches 0; Indels
Sequence 18 BP; 2 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
                             2.8%; S
100.0%;
                 Query Match
Best Local Similarity 100.0
....hes 12; Conservative
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ABZ24285 standard; DNA; 18 BP 335 CGACCAGGGCCG 346 3 ccaccaccicc 14 RESULT 944 ABZ24285/c ଚ

TAA1; wheat; anther; fatty acyl Co-A reductase; FAR; plant; dwarfism; transgenic; lipid metabolism; plant growth; dermatological; octacosanol; fatty alcohol; pharmaceutical; nutritional; dietary; PCR; primer; 88. Wheat TAA1 cDNA RACE antisense primer OL2883.

(first entry)

14-APR-2003

ABZ24285;

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The invention relates to novel isolated and purified polynuclectides, designated TAA1 genes, endogenously expressed in wheat anthers and encode polypeptides having fatty acyl Co-A reductase (FAR) activity. The TAA1 genes are used to produce transgenic plants where the sequence expressed alters lipid metabolism of the transgenic plant. The octacosanol derived from the transgenic plant is useful as a wax, cleaning agent, cosmetic agent, dermatcological agent, pharmaceutical agent, nutritional agent or as a coating agent. A composition comprising a fatty alcohol derived from the transgenic plant is useful in a method of treating or preventing a medical condition. The methods are useful for providing a dietary supplement, the production and isolation of fatty alcohols, and for inducing dwarfism in plants. The methods and other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   compositions of the present invention are useful for the production of transgenic plants and other organisms that comprise increased or altered levels of fatty alcohols used as nutritional or pharmaceutical compositions. Sequences ABZ24280-86 represent primers used for isolating
                                                                                                                                                                                                                                                                                                                       New isolated and purified anther-specific TAA1 nucleotide sequence, useful for the production of transgenic plants with increased or altered levels of fatty alcohols used as nutritional or pharmaceutical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6
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                                                                                                                                                                                                                                          Xia Q, Xie W;
                                                                                                                                                                                                                                                                                                                                                                                                                               Example, Page 46; 124pp; English.
                                                                                                                                                                                                   (CANA ) NAT RES COUNCIL CANADA.
                                                                                                                                                                                                                                             Wang A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    compositions. Sequenc
the wheat TAAl genes
                                                                                                                                                                                                                                                                                    WPI; 2003-167346/16.
                                                                                                                                                                                                                                                                                                                                                                                      compositions.
                                                                                                                                                                                                                                             Selvaraj G,
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Query Match
2.8%; Score 12; DB 1; Length 18;
Best Local Similarity 100.0%; Pred. No. 5.1e+02;
Matches 12; Conservative 0; Mismatches 0; Indels 琚. ADC26391 standard; DNA; 18 337 ACCAGGGCGGG 348 ADC26391; RESULT 945 ADC26391 임 ઠે

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NOV; cytostatic; metabolic disorder; immune; neurodegenerative; circulatory; haemopoletic; wasting; cancer; gene therapy; vaccine; transgenic; human; ss; PCR; primer. NOV protein-related reverse PCR primer SEQ ID 216. 18-DEC-2003

WO2003004687-A2 Homo sapiens.

16-JAN-2003

BP.

(first entry)

97US-00951648. 98US-00174437.

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Identifying a specific binding partner of phosphodiesterase 8 (PDE8) useful for purifying PDE8 products in fluid samples comprises contacting PDE8 with a compound and detecting binding.
                                                                                                             Phosphodiesterase 8; PDE8; human; FB66a; primer; ss.
                                                                                    Human FB66a DNA sequencing primer, W48A9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 3; Col 10; 37pp; English.
                                                                                                                                                                                                                                 11-OCT-2000; 2000US-00686055
AAD59994 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-719642/68.
                                                                                                                                                                                                                                                                                                           (ICOS-) ICOS CORP
                                                                                                                                                                          US6566087-B1.
                                                                                                                                                                                                                                                               16-OCT-1997;
16-OCT-1998;
                                                                                                                                              Homo sapiens
                                                         18-DEC-2003
                                                                                                                                                                                                     20-MAY-2003
                                                                                                                                                                                                                                                                                                                                         Loughney K;
 à
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel isolated NOV polypeptide. The polypeptide of the invention demonstrates cytostatic activity and may be used for determining the presence of, or predisposition to a disease associated with altered levels of expression of the polypeptide, including metabolic disorders, immune disorders, neurodegenerative disorders, circulatory diseases, haemopoietic disorders, wasting diseases and cancer. The polypeptide may also be utilised during gene therapy procedures, vaccine development and transgenic animal production. The current sequence is that of the PCR primer of the invention which was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New isolated NOVX polypeptide, useful for determining the presence of, or predisposition to a disease associated with altered levels of expression of the polypeptide, and for treating or preventing cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gorman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Anderson DW, Berghs C, Boldog FL, Burgess CE, Casman SJ; Catterton E, Edinger S, Eisen AJ, Ellerman K, Gerlach V, GG Guo X, Joffers M, Kekuda R, Li L, Malyankar UM, Miller CE; Padigaru M, Patturajan M, Pena CEA, Rastelli L, Shenoy S; Shimkets RA, Spaderna SK, Spytek KA, Stone DJ, Taupier RJ; Vernet CAM, Voss EZ, Zhong M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 2.8%; Score 12; DB 1; Length 18; Best Local Similarity 100.0%; Pred. No. 5.1e+02; Matches 12; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 18 BP; 4 A; 5 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example C; SEQ ID NO 216; 478pp; English.
                                                                                                                                                                                          17-AUG-2001; 2001US-0313328P.
12-SEP-2001; 2001US-0318711P.
19-SEP-2001; 2001US-032338P.
21-SEP-2001; 2001US-0323969P.
04-JAN-2002; 2002US-0345022P.
04-JAN-2002; 2002US-0345038P.
28-FEB-2002; 2002US-0345038P.
                             05-JUL-2001; 2001US-0303046P.
09-JUL-2001; 2001US-0303828P.
09-JUL-2001; 2001US-0304016P.
11-JUL-2001; 2001US-0304528P.
13-JUL-2001; 2001US-0305262P.
17-JUL-2001; 2001US-0305673P.
17-JUL-2001; 2001US-030685P.
24-JUL-2001; 2001US-030828P.
30-JUL-2001; 2001US-0309258P.
17-AUG-2001; 2001US-0309258P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                used to analyse human NOV DNA.
                                                                                                                                                                                                                                                                                                                                                                             16-APR-2002; 2002US-0372326P.
16-APR-2002; 2002US-0372390P.
19-APR-2002; 2002US-0373881P.
19-APR-2002; 2002US-0373881P.
                                                                                                                                                                                                                                                                                                2002US-0360814P.
2002US-0360830P.
2002US-0361133P.
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                                                                                                                                                                                                                                                                                                                                                                           02US-0363637P
                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-JUL-2002; 2002US-00188186
      03-JUL-2002; 2002WO-US021361
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-221607/21.
                                                                                                                                                                                                                                                                                                                                             01-MAR-2002;
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The invention relates to a method for identifying a specific binding partner of phosphodiesterase 8 (PDE8). The method is useful for identifying a specific binding partner of PDE8, which inhibits or enhances activity of PDE8. The binding partners of PDE8 are useful for purification, detection or quantification of PDE8 products in fluid and tissue samples using immunological procedures. Modulators of PDE8 activity are useful in treating a wide range of diseases and physiological conditions in which PDE8 activity is known to be involved. The present sequence is a primer used for sequencing human PDE8 AZ splice variant DNA (FB66a)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Complementarity-determining region; CDR; humanised; antibody; hIL2R; human; interleukin; IL-2; receptor; murine; anti-human; Ab; T-cell; monoclonal antibody; B-B10; mixed lymphocyte reaction; variable; V; region; PCR; framework; plamid; heavy; H; light; L; amplify; primer; polymerase chain reaction; 88.
                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                      ;
0
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2.8%; Score 12; DB 1; Length 18;
Best Local Similarity 100.0%; Pred. No. 5.1e+02;
Matches 12; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                              Sequence 18 BP; 4 A; 7 C; 2 G; 5 T; 0 U; 0 Other;
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AAQ43232 standard; DNA; 15 BP.
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13-OCT-1993 (first entry)
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Gaps

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344 CCGGCTGCTCTA 355 CCGGCTGCTCTA 13

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RESULT 946 AAD59994

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(BIOG-) BIOGNOSTIK GES BIOMOLEKULARE DIAGNOSTIK.

Brysch W;

Schlingensiepen K,

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The sequences given in AAQ43226-32 are primers which were used in the cloning of DNA encoding the variable (V) regions of the murine anticoloning of DNA encoding the variable (V) regions of the murine anticoloning of DNA encoding the variable (V) regions of the murine anticoloning the variable (V) This MAD was used in the construction of a humanised antibody (Ab) which binds specifically to human interleukin (IL)-2 receptor (MIL2R). The complementarity-construction of a human form of the house of the House of the House AR37599-04). The DNA encoding the variable (V) region of IL-2 to the IL-2 receptor on human T-cells. It also inhibits the human mixed lymphocyte reaction. The DNA encoding the variable (V) region of the B-10 Ab was cloned by PCR and sequence homology to the murine sequence was selected and the framework of this Ab was bound with the B-B10 V was selected and the framework of this Ab was bound with the B-B10 V region CDR and a part of the framework to design several kinds of the humanised B-B10 V region. The DNA sequence coding this humanised B-B10 was constructed.
                                                                                                                                                                                                                                                                                        Humanised antibody comprising - CDR region of mouse MAB B-B10 specific for IL-2 receptor useful for treating carcinoma expressing IL-2 receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   c-fos; antisense oligonucleotide; modulate; gene expression; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cuery Match
2.8%; Score 11.8; DB 1; Length 15;
Best Local Similarity 86.7%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15 BP; 1 A; 7 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 c-fos gene antisense oligonucleotide c-fos-22.
                                                                                                                                                                                                                          Wijdenes J, Noguchi H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Updated on 25-MAR-2003 to correct PN field.
                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 45; 62pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAV48908 standard; DNA; 15 BP
                                                                                                                                                    SUMU ) SUMITOMO PHARM CO LTD.
BIOT ) BIOTEST PHARMA GMBH.
INNO-) INNOTHERAPIE LAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97EP-00101531
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      282 GGCACCAAGCTGGTG 296
                                                                                   92WO-JP001583
                                                                                                                   91JP-00323319
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                                                                                                                                                                                                                          Nakatani T, Gomi H,
                                                                                                                                                                                                                                                            WPI; 1993-197057/24.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-JAN-1997;
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                                                                                   03-DEC-1992;
                                                                                                                      16-DEC-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EP856579-A1
                  WO9311238-A1
                                                  10-JUN-1993.
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AAV48887-929 represent antisense oligonucleotides directed against the c-
fos gene. Of these, only oligonucleotides AAV48887-917 resulted in
c fos gene. Of these, only oligonucleotides AAV48887-917 resulted in
c significant redocution in c-fos protein expression, while oligonucleotides
cc AAV48918-29 had little effect. The oligonucleotides exemplify the
contain or specification describes oligonucleotides that contain 8-30
c invention. The specification describes oligonucleotides that contain there
cc nucleotides, which contain at most 8 nucleotides that can each form three
cc ontain two sequences of three consecutive nucleotides each able to form
cc contain two sequences of three consecutive nucleotides each able to form
cc contain two sequences of three consecutive nucleotides each able to form
cc contain two sequences of three consecutive nucleotides each able to form
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cc contain two sequences of the oligonucleotides are used to one
cc given by 2R/3R = 0.33-0.72. The oligonucleotides are used to one odulate
cc given by 2R/3R = 0.30-0.72. The oligonucleotides can also be used to analyse
cc and/or keratinocytes). The oligonucleotides can also be used to analyse
cc therapeutically, e.g. in cases of cancer or (targeting TGF) for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                              Preparation of antisense oligo:nuclectide(s) which lack long runs of consecutive guanosine or inosine - and have specific ratio of residues able to form two or three hydrogen bonds, have greater activity and reduced toxicity, used therapeutically or to modulate growth of cells in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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86.7%; Pred. No. 3.8e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 15 BP; 2 A; 4 C; 5 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 10; Fig 7; 286pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             stimulating the immune system
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Best Local Similarity 86.7'
Matches 13; Conservative
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                                                                                                                                                WPI; 1998-400910/35
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Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31-JAN-1997;
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AAV48887-929 represent antisense oligomucleotides directed against the cfos gene. Of these, only oligomucleotides AAV48887-917 resulted in
significant redcution in c-fos protein expression, while oligomucleotides
AAV48918-29 had little effect. The oligomucleotides exemplify the
incleotides, which contain at most 8 nucleotides that contain 8-30
nucleotides, which contain at most 8 nucleotides that contain 8-30
nucleotides, which contain at most 8 nucleotides that contain 8-30
contain two sequences of three consecutive cytosines; do not
contain two sequences of three consecutive ortosides each able to form
three H-bonds each to four consecutive cytosines; do not
contain two sequences of three consecutive ortosides each able to form
three H-bonds to three consecutive ortosines, and the ratio between
creaidues able to form two H-bonds each (2R) or three such bonds (3R) is
creaidues able to form two H-bonds each (2R) or three such bonds (3R) is
creationed as a particularly the genes for p53, EB-2, june, juno,
creation bonds are an ilver or kidney cells, osteoclasts, osteoblasts
candyor keratinocytes). The oligomucleotides can also be used to analyse
thuction of proteins (by altering their expression or activity) and
the transpeutically, e.g. in cases of cancer or (targeting TGF) for Preparation of antisense oligo:nucleotide(s) which lack long runs of consecutive guanosine or inosine - and have specific ratio of residues able to form two or three hydrogen bonds, have greater activity and reduced toxicity, used therapeutically or to modulate growth of cells in claim 10; Fig 7; 286pp; English. culture

Sequence 15 BP; 4 A; 3 C; 6 G; 2 T; 0 U; 0 Other;

ö Gaps . Query Match
2.8%; Score 11.8; DB 1; Length 15;
Best Local Similarity 86.7%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 2; Indels

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AAV48699 standard; DNA; 15 15-OCT-1998 AAV48699; 950 임

BP.

junB gene antisense oligonucleotide JunB-T-8. (first entry)

junB; junD; antisense oligonucleotide; modulate; gene expression; ss.

Homo sapiens

Synthetic

05-AUG-1998 EP856579-A1

97EP-00101531 31-JAN-1997; (BIOG-) BIOGNOSTIK GES BIOMOLEKULARE DIAGNOSTIK.

97EP-00101531

31-JAN-1997;

Brysch W; Schlingensiepen K,

WPI; 1998-400910/35.

Preparation of antisense oligo:nucleotide(s) which lack long runs of consecutive guanosine or inosine - and have specific ratio of residues able to form two or three hydrogen bonds, have greater activity and reduced toxicity, used therapeutically or to modulate growth of cells in

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AAV48564-708 represent antisense oligonucleotides directed against the cube and junb genes. Of these, only oligonucleotides AAV4855-614

Tresulted in effective downregulation of negative growth control by Junb or Junb, while AAV48615-708 had little effect. The oligonucleotides exemplify the invention. The specification describes oligonucleotides compared to the contain 8-30 nucleotides, which contain a most B nucleotides that can each form three hydrogen bonds to cytosine; do not contain four consecutive mucleotides able to form three H-bonds each to four consecutive cytosines; do not contain two sequences of three consecutive cycosines, and the ratio between residues able to form two H-bonds each cytosines, and the ratio between residues able to form two H-bonds each cytosines, and the ratio between residues able to form two H-bonds each cytosines, and the ratio between residues able to form two H-bonds each cytosines, and the ratio between residues able to form two H-bonds each cytosines, and the ratio between residues able to form two H-bonds each cytosines, and the rased to modulate expression of genes, particularly configuration of primary cell cultures (e.g. bone marrow stem, liver or kidney cells, osteoclasts, ost
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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2.8%; Score 11.8; DB 1; Length 15;
Best Local Similarity 86.7%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 2; Indels
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                                                                          Example 3; Fig 5c; 286pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     147 GTGGAGGCCGGCTTC 161
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AAX31429 standard; DNA; 15 BP. RESULT 951 AAX31429

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21-MAY-1999 (first entry) AAX31429;

Tag sequence; colorectal cancer; pancreatic cancer; colon cancer; diagnosis; prognosis; treatment; ss. Tag sequence of a transcript decreased in colorectal cancer.

Homo sapiens

WO9853319-A2

26-NOV-1998.

98WO-US010277. 20-MAY-1998; 21-MAY-1997;

(UYJO) UNIV JOHNS HOPKINS. Vogelstein B, Kinzler KW;

WPI; 1999-070161/06.

Use of isolated gene transcripts - useful for developing products for the diagnosis, prognosis and treatment of cancers, particularly colon and pancreatic cancer.

Claim 1, Page 50, 120pp; English.

AAX10947-31815 represent tag sequences of transcripts that are differentially expressed in colorectal cancer, in pancreatic cancer, in both. The tag sequences can be used to identify genes by matching

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Gaps

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AAX30947-31815 represent tag sequences of transcripts that are differentially expressed in colorectal cancer, in pancreatic cancer, or in both. The tag sequences can be used to identify genes by matching the tag to a gen date base member, or by using the tag sequences as probes to isolate unidentified genes from cDNA libraries. The tag sequences can slow be used in a method for diagnoshing colon or pancreatic cancer in a sample suspected of being neoplastic. The method comparises comparing the level of at least one transcript in a first sample of a tissue to a second sample, where the first sample is a colonic tissue suspected of being neoplastic and the second sample is a normal human colonic tissue. The transcript is identified by a tag selected from AAX30947-31815. The methods of the invention can be used in the diagnosis, prognosis and
tag to a gen data base member, or by using the tag sequences as probes to also be used in a method for diagnosing rollon or pancratic cancer in a sample suspected of being neoplastic. The method comprises comparing the level of at least one transcript in a first sample of a tissue to a second sample, where the first sample is a colonic tissue suspected of being neoplastic and the second sample is a normal human colonic tissue methods of the invention can be used in the diagnosis, prognosis and treatment of cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Use of isolated gene transcripts - useful for developing products for the diagnosis, prognosis and treatment of cancers, particularly colon and pancreatic cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tag sequence; colorectal cancer; pancreatic cancer; colon cancer; diagnosis; prognosis; treatment; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tag sequence of a transcript increased in pancreatic cancer.
                                                                                                                                                                                                                                                                  Query Match 2.8%; Score 11.8; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                             Sequence 15 BP; 4 A; 1 C; 6 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX31675 standard; DNA; 15 BP
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                                                                                                                                                                                                                                                                                                                                                   287 CAAGCTGGTGAAGGA 301
                                                                                                                                                                                                                                                                                                                                                                                           1 CATGTTGGTGAAGGA 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-070161/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-MAY-1998;
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Sequence 15 BP; 4 A; 1 C; 6 G; 4 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX59550-53 represent intron/exon and exon/intron junctions of the mouse pitx3 gene. Pitx3 proteins are homeobox domain proteins, which are involved in the development of the lens and contribute to diseases and disorders of the lens, such as cataracts. The Pitx3 nucleic acids (e.g. antisense sequences, ribozymes and triplex nucleic acids), probes derived from them and polypeptides, are useful in claimed methods to detect an ocular disease, especially of the lens, e.g. cataract formation. Specific conditions that can be detected and treated are Anterior Segment
                                   Gaps
                                                                                                                                                                                                                                                                                                                          Pitx1; homeobox domain protein; lens development; lens disorder; cataract; detection; ocular disease; ASMD; Peter's anomaly; anterior segment mesenchymal dysgenesis; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pitx3, homeobox protein, and related nucleic acid sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.8%; Score 11.8; DB 1; Length 15;
86.7%; Pred. No. 3.8e+02;
tive 0; Mismatches 2; Indels
   Length 15;
                                   2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Forward primer #78 used in multiplexing PCR/SBE assay.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mesenchymal Dysgenesis (ASMD) and Peter's anomaly
Query Match 2.8%; Score 11.8; DB 1; Best Local Similarity 86.7%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                            Intron 2/exon 3 juntion of the mouse Pitx3 gene.
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                                                                                                                                                                                       AAX59553 standard; DNA; 15 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97US-00957351.
                                                                  287 CAAGCTGGTGAAGGA 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (IOWA ) UNIV IOWA RES FOUND.
                                                                                                  1 cargricercaacea 15
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                                                                                                                                                                                                                                                         21-JUL-1999 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Semina EV, Murray JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1999-312965/26.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9921996-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-OCT-1998;
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                                                                                                                                                                                                                         AAX59553;
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                                                                                                                                                                       AAX59553,
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Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or

Example 7; Page 47; 201pp; English.

inflammation.

Edmondson SR;

Werther GA,

Wraight CJ,

WPI; 2001-041421/05.

(MURD-) MURDOCH CHILDRENS RES INST.

21-JUN-2000; 2000WO-AU000693

28-DEC-2000.

21-JUN-1999;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to an oligonucleotide array comprising array bigloanucleotide tags fixed to a solid substrate. The oligonucleotide array is useful for genotyping a nucleic acid sample at one or more loci via single base extension (SBE) reactions. A pair of primers is used to amplify a polymorphic locus in a sample e.g. a single nucleotide polymorphism (SNP). The present sequence is one of the primers used in the method of the present invention to amplify a polymorphic sample. The amplification uncleotide product is then used as a template in a SBE reaction with an extension primer. The SBE reaction products are used to form the oligonucleotide array
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.8%; Score 11.8; DB 1; Length 15;
86.7%; Pred. No. 3.8e+02;
Ative 0; Mismatches 2; Indels 0; Gaps
                           Oligonucleotide array; genotyping; single base extension reaction, SBE; PCR primer; polymorphic locus; single nucleotide polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Universal array of oligonucleotides tags attached to a solid substrate along with locus-specific tagged oligonucleotides useful in genotyping using single base extension reactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hirschhorn JN, Huang X, Kaplan P, Lander ES, Lockhart DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15 BP; 4 A; 5 C; 6 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (WHED ) WHITEHEAD INST BIOMEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 7; Page 56; 70pp; English.
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                                                                                                                                                                                                                                                                                                                                                               27-MAR-2000; 2000WO-US008069.
                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-0126473P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (AFFY-) AFFYMETRIX INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-656171/63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ryder T, Sklar P;
                                                                                                                                                                                                             WO200058516-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-JUN-1999;
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                                                                                                                                          Unidentified
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ID AAF4

AC AAF4

DE AAF4

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The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an anticomplete contacting the skin with an anticomplete contacting the skin with an receptor, IGF binding protein [GFEPP]-2 or IGFEPB3), which is capable of inhibiting or reducing growth factor mediated cell proliferation. Inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153 or P45161). The method is useful for ameliorating the effects of psoriasis, rochthyosis, pityriasis, ruba, plaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperrecovascular condition such as a neovascular condition of the retina, branch service with a prowth factor mediated malignancies, other sclerotic disease, kidney disease, hyperplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antisense therapy, antiproliferative, antinflammatory, antipsoriatic, cytostatic, dermatological, cardiant, virucide, ophthalmological, keloid; skin disorder, insulin-like Growth Factor I receptor; IGP-1; pityriasis; IGF binding protein, IGFBP-2; IGFBP3; inflammation, psoriasis; pilatis; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba, keratosis; neophasia; scleroderma; wart, skin cancer; sclerotic disease; hypermeovascular condition; hyperplasia; kidney disease; neobvascular condition of the retina; ss.
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WO200078341-A1

Homo sapiens

Edmondson SR;

Werther GA,

Wraight CJ,

Edmondson SR;

99US-0140345P

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Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin discorder; Insulin-like Growth Factor. I receptor; IGF1; pityriasis; IGF binding procein; IGFBP-2; IGFBP3; inflammation; psoriasis; prowth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neophasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplama; kidney disease; neobarchoo of the retina; sidney disease;
                                                                                                                                                                                                                                                                                                        The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonuclectide, (for Insulin-IRe Growth Factor [IGF].

receptor, IGF binding protein [IGFBP] - 2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonuclectide which can be used to design the antisense oligonuclectide which can be used to design the antisense oligonuclectides of the present invention (see AAF4151 and AAF45153 and AF45163), ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hypernevascular condition and as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood
                                                                                                                                                                        Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 2.8%; Score 11.8; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 15 BP; 2 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                         (MURD-) MURDOCH CHILDRENS RES INST.
                                                                                                                                                                                                                                                                            Example 8; Page 72; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      vessels or any other hyperplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF47144 Standard; DNA; 15 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               147 GTGGAGGCCGGCTTC 161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IGFBP3 oligonucleotide #564.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 Grecadecedecare 15
                                                                                             Wraight CJ, Werther GA,
                                                                                                                                    WPI; 2001-041421/05
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                                                                                                                                                                                                                                          .nflammation.
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                   21-JUN-1999;
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(first entry)

(MURD-) MURDOCH CHILDRENS RES INST.

99US-0140345P.

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The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligomucleotide, (for Insulin-like Growth Factor [IGR]-1. Comprises contacting the skin with an acceptor, IGF binding protein [IGRP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an inflammation and/or other disorders. The present sequence is an oligomucleotide which can be used to design the antisense oligomucleotide which can be used to design the antisense oligomucleotide of the present invention (see AAP45151 and AAF45153-0190nucleotides of the present invention (see AAP45151 and AAF45153-0190nucleotides of the present invention growths, cancers of psoriasis, rethynosis, playing growths, cancers of the skin, a neoplasias, scleroderma, warts, benign growths, cancers of the skin, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antisense therapy; antiproliferative; antinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; sein discorder; Insulin-like Growth Factor I receptor; IGF-1; pityrisais; IGF binding protein; IGFB-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keatosis; neophasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neobascular condition; hyperplasia; kidney disease;
                                                                               Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.8%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 3.8e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 15 BP; 2 A; 7 C; 3 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                        Example 7; Page 47; 201pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   322 TGCTGGCGGCACG 336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IGF-I oligonucleotide #1302.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAF50342 standard; DNA; 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 2.8
Best Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-041421/05
                                                      WPI; 2001-041421/05
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Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                inflammation.
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Example 8; Page 69; 201pp; English.

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an anticontaction of the skin with an anticontaction and specific (for Insulin-like Growth Factor [168]-1 receptor, IdF binding protein [168P]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, or cher disorders. The present sequence is an inflammation and/or other disorders. The present sequence is an inflammation and/or cher disorders. The present sequence is an inflammation and/or the present invention (see AAF45151 and AAF45153-1) coligonucleotide which can be used to design the antisense of the present invention (see AAF45151 and AAF45153-1) coligonucleotide of the present invention (see AAF45151 and AAF45153-1) coligonucleotides of the present invention (see AAF45151 and AAF45153-1) coligonucleotides of the present invention growths, cancers of the skin, a necessary proper section of the skin, a brain or skin, growth factor-mediated malignancies on of the retina, the present of the section of the section disease, kidney disease, hyperplasia

Sequence 15 BP; 2 A; 6 C; 2 G; 5 T; 0 U; 0 Other;

ö Gaps ö 2.8%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 3.8e+02; ive 0; Mismatches 2; Indels 13; Conservative Similarity Query Match Best Local 9 Matches

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RESULT 959 AAF47290

AAF47290 standard; DNA; 15 BP AAF47290;

IGFBP3 oligonucleotide #710.

(first entry)

30-MAR-2001

Antisense therapy, antiproliferative, antiinflammatory, antipsoriatic, cytostatic, dermatological; cardiant; virucide; ophthalmological; keloid; skin discorder; insulin-like Growth Factor 1 receptor; IGFP-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; plantis; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hypermeovascular condition; hyperplasia; kidney disease; neobascular condition; hyperplasia; kidney disease;

Homo sapiens.

WO200078341-A1.

28-DEC-2000

21-JUN-2000; 2000WO-AU000693

MURD-) MURDOCH CHILDRENS RES INST. 21-NUT-1999;

99US-0140345P.

Werther GA, Edmondson SR; Wraight CJ,

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or

inflammation

Example 7; Page 48; 201pp; English.

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an stin disorders. The method comprises contacting the skin with an capturent of the property of the state of state

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Sequence 15 BP; 3 A; 6 C; 3 G; 3 T; 0 U; 0 Other;

Gaps ò Query Match 2.8%; Score 11.8; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 2; Indels 13; Conservative Matches

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AAF52600 standard, DNA; 15 RESULT 960

AAF52600;

(first entry) 30-MAR-2001

IGF-I oligonucleotide #3560.

Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; selvi discorder; insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFB-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neophasia; scleroderma; wart; skin cancer; sclerotic disease; hyperaneovascular condition; hyperplasia; kidney disease; neovascular condition; hyperplasia; kidney disease;

Homo sapiens,

WO200078341-A1.

28-DEC-2000.

21-JUN-2000; 2000WO-AU000693.

21-JUN-1999;

(MURD-) MURDOCH CHILDRENS RES INST.

Wraight CJ, Werther GA, Edmondson SR;

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation

Example 8; Page 84; 201pp; English.

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonuclectide, (for Insulin-like Growth Factor [IGRF-1] and sector [IGRF-2]. The present sequence is an inflammation and/or other disorders. The present sequence is an oligonuclectide which can be used to design the antisense oligonuclectide which can be used to design the antisense of the present invention (see AAF45151 and AAF45153-CFF45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborthoea, keloids, keratosis, chhyperneovascular condition such as a neovascular condition of the skin, a hyperneovascular condition such as a neovascular condition chief service disease, kidney disease, hyperproliferation of the inside of blood classas, custom the hyperplasia

Seguence 15 BP; 0 A; 6 C; 6 G; 3 T; 0 U; 0 Other;

Gaps ; 0 2.8%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 3.8e+02; tive 0; Mismatches 2; Indels Local Similarity 86.7 Query Match

GGCTGCTTCCCGGGC 254 1 decrectected 240

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AAF47145/c ID AAF47145 standard; DNA; 15 RESULT 961

AAF47145;

GFBP3 oligonucleotide #565

(first entry)

30-MAR-2001

Antisense therapy; antiproliferative; antinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; gath discorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding procein; IGFB-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neophasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasis, kidney disease; neobascular condition; hyperplasis, kidney disease;

Homo sapiens.

WO200078341-A1.

21-JUN-2000; 2000WO-AU000693 99US-0140345P 21-JUN-1999;

(MURD-) MURDOCH CHILDRENS RES INST.

Edmondson SR; Wraight CJ, Werther GA,

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or nflammation.

Example 7; Page 47; 201pp; English.

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonuclectide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of

infibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense April 1900 of Jugonucleotides of the present invention (see AAF45151 and AAF45153 of the method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood vessels or any other hyperplasia

889999999998888

Sequence 15 BP; 2 A; 7 C; 3 G; 3 T; 0 U; 0 Other;

Gaps ö 2.8%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 3.8e+02; ive 0; Mismatches 2; Indels Matches 13; Conservative Query Match Best Local Similarity

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321 GIGCIGGCGGCGAC 335 15 Grecresadaceak 1 셤 ઠે

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RESULT 962 AAF45774

AAF45774 standard; DNA; 15 BP.

AAF45774;

(first entry) 30-MAR-2001 IGFBP2 oligonucleotide #613.

Antisense therapy; antiproliferative; antinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disporder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBPB; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis, neophasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasis, kidney disease; neobascular condition; hyperplasis, kidney disease;

Homo sapiens.

WO200078341-A1

28-DEC-2000.

21-JUN-2000; 2000WO-AU000693.

99US-0140345P. 21-JUN-1999; (MURD-) MURDOCH CHILDRENS RES INST.

Edmondson SR;

Werther GA,

Wraight CJ,

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or

Example 6; Page 38; 201pp; English.

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antiesnee oligonucleotide, (for Insuln-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFSP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, infilammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153-

F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor—mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood vessels or any other hyperplasia 88666666888

Sequence 15 BP; 4 A; 4 C; 5 G; 2 T; 0 U; 0 Other;

Query Match

2.8%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 3.8e+02; ative 0; Mismatches 2; Indels GIGGACATCACCACG 101 Best Local Similarity 86.7 Matches 13; Conservative δ

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1 Grecacacacard 15

AAF46991 standard; DNA; 15 BP AAF46991; RESULT 963 **AAF4**699

IGFBP3 oligonucleotide #411 30-MAR-2001 (first entry)

Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; spin discorder; insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding procein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilatis; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keatoolsis, neophasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition; hyperplasia; kidney disease;

Homo sapiens.

WO200078341-A1.

21-JUN-2000; 2000WO-AU000693

21-JUN-1999; 99US-0140345P.

(MURD-) MURDOCH CHILDRENS RES INST.

Edmondson SR; Wraight CJ, Werther GA,

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.

Example 7; Page 46; 201pp; English.

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF] receptor, IGF binding protein [IGFBP] 2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153 + F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, ruba, pilaris, serbornhoea, keloids, keratosis, neoplasis, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina,

disease, kidney disease, hyperproliferation of the inside of blood vessels or any other hyperplasia ឧឧឧ

Sequence 15 BP; 2 A; 6 C; 6 G; 1 T; 0 U; 0 Other;

ô 2.8%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 3.8e+02; tive 0; Mismatches 2; Indels Query Match Best Local Similarity 86.7 Matches 13; Conservative

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108 CGCGACCGCAGCAAG 122 a ò

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Gaps .. 0

BP RESULT 964 AAF47146/c ID AAF47146 standard; DNA; 15

AAF47146;

(first entry) 30-MAR-2001 IGFBP3 oligonucleotide #566.

Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disporder; insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFB-2; IGFBP3; inflammation; psoriasis; plaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; soleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasis; kidney disease; neobascular condition; hyperplasis; kidney disease;

Homo sapiens.

WO200078341-A1.

28-DEC-2000.

21-JUN-2000; 2000WO-AU000693.

99US-0140345P. 21-JUN-1999; (MURD-) MURDOCH CHILDRENS RES INST.

Edmondson SR;

Werther GA,

Wraight CJ,

WPI; 2001-041421/05

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.

Example 7; Page 47; 201pp; English

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonuclectide, (for Insulin-like Growth Factor [IGFP]-1 creceptor, IGFP binding protein [IGFPP]-2 or IGFPP), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonuclectide which can be used to design the antisense oligonuclectide of the present invention (see AAF45151 and AAF45153-0190uclectides of the method is useful for ameliorating the effects of psoriasis, not invention such as a neovascular condition of the retina, hyperneovascular condition such as a neovascular condition of the retina, disease, hyperproliferation of the inside of blood disease, hyperproliferation of the inside of blood vessels or any other hyperplasia

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The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an anticense oligonucleoride, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleoride which can be used to design the antisense APF45153 and APF45153 or ligonucleorides of the present invention (see APF45151 and APF45153 or P45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pliaris, serborthoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a neoplasias, stun, growth factor-mediated malignandses, other sclerotic brain or skin, growth factor-mediated malignandses, other sclerotic conserved.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                              Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; shi discorder; insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; plaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; kearbosis; neoplasia; geleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasis; kidney disease; neoblasis; schorker condition; hyperplasis; kidney disease;
                                                                            Gaps
                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 11.8; DB 1; Length 15; Pred. No. 3.8e+02;
                                    Length 15;
                                                                          2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seguence 15 BP; 5 A; 6 C; 3 G; 1 T; 0 U; 0 Other;
Sequence 15 BP; 2 A; 7 C; 3 G; 3 T; 0 U; 0 Other;
                                    Score 11.8; DB 1;
Pred. No. 3.8e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Werther GA, Edmondson SR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MURD-) MURDOCH CHILDRENS RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 8; Page 75; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ressels or any other hyperplasia
                                                                                                                                                                                                                                                    AAF51317 standard; DNA; 15 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-JUN-2000; 2000WO-AU000693.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-0140345P.
                                  2.8%;
                                                                                                                   320 CGTGCTGGCGGCGGA 334
                                                                                                                                                                                                                                                                                                                                                                 IGF-I oligonucleotide #2277.
                                                                                                                                                     cerecreeaeacea 1
                                                                                                                                                                                                                                                                                                                               (first entry)
                                    2.87
Best Local Similarity 86.77
Matches 13, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-041421/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200078341-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wraight CJ,
                                                                                                                                                                                                                                                                                                                               30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-DEC-2000.
                                                                                                                                                                                                                                                                                           AAF51317;
                                                                                                                                                         13
                                                                                                                                                                                                                  RESULT 965
                                                                                                                                                                                                                                    AAF51317
                                                                                                                                                                                                                                                                        g
                                                                                                                       δ
                                                                                                                                                     q
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2.8%;

Query Match Best Local Similarity

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The present invention describes genetic variants of the human glutathione ceductase (GSR) gene (1). (1) has antianaemic activity and can be used in reductase (GSR) gene (1). (1) has antianaemic activity and can be used in screening for drugs teargeting (1) that come the used in screening for drugs teargeting (1) that come the present come to be used; for improving the efficiency and reliability of several steps in the discovery and development of drugs for treating diseases associated with GSR activity; for haplotyping, which is also used by the pharmaceutical research scientist to validate GSR as a candidate target for treating a specific condition of disease predicted to be associated with GSR activity, end for screening compounds targeting GSR. Company of the capturest of the service of the variation of the binding affinity of candidate drugs to treat the binding affinity of candidate drugs to treat come company are variation on the biological activity of GSR activity. (1) is also useful in studying the captesent of the variation of the binding affinity of candidate drugs targeting GSR for the treatment company affinity of candidate drugs targeting GSR for the treatment company. The present sequence represents an allele specific company as single nucleotide polymorphism) in the ASO primer is shown cusing an IUPAC ambiguity code (as given in the present invention)
ô
                                                                                                                                                                                                                                                                                            Human; glutathione reductase; GSR; enzyme; haemolytic anaemia; SNP;
gene therapy; antianaemic; polymorphic; single nucleotide polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New genetic variants of Glutathione reductase isogenes, useful for improving efficiency and reliability in drug development for treating hemolytic anemia.
 Gaps
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                                                                                                                                                                                                                                                           Human GSR allele specific oligonucleotide primer SEQ ID NO:34.
   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15 BP; 1 A; 9 C; 4 G; 0 T; 0 U; 1 Other;
   5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sun X;
     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                           /*tag= a
/note= "polymorphic base"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sanchis A, Sausker EA,
                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 14; Page 14; 137pp; English
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GENA-) GENAISSANCE PHARM INC.
                                                                                                                                                           ABN87915 standard; DNA; 15 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-NOV-2000; 2000US-0247202P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-NOV-2001; 2001WO-US046473
                                      86 AGTGGACATCACCAC 100
                                                                        15
                                                                                                                                                                                                                               (first entry)
     13; Conservative
                                                                    1 AGTGGCCAACACCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-471719/50.
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                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                        misc_feature
                                                                                                                                                                                                                               12-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-MAY-2002
                                                                                                                                                                                                                                                                                                                                           primer; ss
                                                                                                                                                                                                ABN87915;
                                                                                                                             RESULT 966
         Matches
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13;

Matches

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15

RESULT 967

ABK32383

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Human; cholinergic receptor muscarinic 5; CHRMS; genotyping; haplotyping;
                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to an isolated, purified human nucleic acid (I) that has the same sequence as a mRNA found in humans and is a SAGE (serial analysis of gene expression) tag comprising a single stranded probe containing at least 10 consecutive nucleotides. SAGE tags, are diagnostic and prognostic markers of cancer, especially of the colon and pancreas. ABK31900-ABK32770 represent human colon and pancreatic cancer SAGE tags of the invention
Human, colon cancer, colorectal cancer, pancreatic cancer; SAGE tag; serial analysis of gene expression, diagnostic, prognostic, probe;
                                                                                                                                                                                                                                                                                                                                            New human nucleic acid containing specific SAGE tags, useful as diagnostic markers for cancer, also derived probes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Denton RR, Nandabalan K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.8%; Score 11.8; DB 1; Length 15; llarity 86.7%; Pred. No. 3.8e+02; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human CHRM5 gene polymorphism detection ASO primer #8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 15 BP; 4 A; 1 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                              3
                                                                                                                                                                                                                                                                              Zhou
                                                                                                                                                                                                                                                                              Zhang L,
                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Col 83; 161pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Choi JY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GENA-) GENAISSANCE PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .782/c
ABK81782 standard; DNA; 15 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-OCT-2001; 2001WO-US032022.
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                                                                                                                                                                             98US-00081646.
                                                                                                                                                                                                              98US-00081646.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CATGTTGGTGAAGGA 15
                                                                                                                                                                                                                                                                                Vogelstein B, Kinzler KW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-AUG-2002 (first entry)
                                                                                                                                                                                                                                               SNING OLYJO ( OLYJ)
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Sausker EA, Stephens JC;
                                                                                                                                                                                                                                                                                                             WPI; 2002-153821/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
Matches 13; Conserv
                                          cancer marker; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer; ss
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                                                                            Homo gapiens
                                                                                                                                                                             20-MAY-1998;
                                                                                                           US6333152-B1
                                                                                                                                                                                                                20-MAY-1998;
                                                                                                                                            25-DEC-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                             Gaps
                                                                                                                                                                                                                                                                                                                                       Human, colon cancer, colorectal cancer, pancreatic cancer, SAGE tag, serial analysis of gene expression; diagnostic, prognostic, probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New human nucleic acid containing specific SAGE tags, useful as diagnostic markers for cancer, also derived probes.
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          2.8%; Score 11.8; DB 1; Length 15;
86.7%; Pred. No. 3.8e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seguence 15 BP; 4 A; 1 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Zhou W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human pancreatic cancer SAGE tag #181.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Zhang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Col 55; 161pp; English
                                                                                                                                                                                                                                                                                                       Human colon cancer SAGE tag #484
                                                                                                                                                                                                     BP
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                                                                                321 GTGCTGGCGGCGAC 335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABK32629 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vogelstein B, Kinzler KW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SNINGO NINU ( OCYU)
                                                                                                                                                                                                       ABK32383 standard; DNA; 15
                                                                                                                 GRGCTGGCGGCGGC 1
                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13; Conservative
                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-153821/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                               cancer marker; ss.
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Homo sapiens US6333152-B1

23-APR-2002

ABK32383;

20-MAY-1998; 20-MAY-1998;

23-APR-2002

ABK32629;

ABK32629
ID ABK3
XX
AC ABK3
XX
DT 23-A
XX

RESULT 968

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Best Loca Matches

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Gaps

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WPI; 2002-435523/46.

Claim 14; Page 13; 72pp; English.

Novel cholinergic receptor, muscarinic 5 polynucleotide useful therapeutically and in screening for candidate drug to treat diseases related to the receptor activity.

The present invention relates to a new cholinergic receptor, muscarinic 5 (CHRMS) polymuclectide comprising a sequence which is a polymorphic variant for a reference sequence for the CHRMS gene or its fragment, or a polymorphic variant of a reference sequence for a CHRMS conA or its fragment. The invention is useful in drug screening assays. The molecules of the invention are useful in studying the expression and function of CHRMS, and in expressing CHRMS protein for use in screening for candidate drugs to treat diseases related to CHRMS activity. The methods of the invention are useful in developing diagnostic tests and therapeutic treatments. The method is also useful in the design of clinical trials of andidate drugs for treating specific condition or disease associated with CHRMS activity and is useful in determining whether an individual has one of the haplotypes or one of the haplotype pairs. The invention is useful in a variety of diagnostic and prognostic formats and therapeutic methods. The invention is also useful in genotyping and/or haplotyping the CHRMS gene in an individual. The present nucleic acid sequence the CHRMS gene in an individual were used in the invention to detect polymorphisms in the human CHRMS gene

Sequence 15 BP; 2 A; 6 C; 4 G; 2 T; 0 U; 1 Other;

Query Match 2.8%; Score 11.8; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 2; Indels ઠે

0; Gaps

RESULT 970

ABZ76557

ABZ76557 standard; DNA; 15 BP.

ABZ76557;

29-APR-2003 (first entry)

Lactobacillus brevis PCR primer ORF4 SEQ ID NO:60.

Lactobacillus brevis; beer turbidity; beer clouding; beer; detection; lactic acid bacteria; brewing; probe; PCR primer; ss.

Lactobacillus brevis.

WO200295028-A1.

28-NOV-2002.

23-MAY-2002; 2002WO-JP005022.

23-MAY-2001; 2001JP-00154085.

(KIRI) KIRIN BEER KK

Fujii T;

WPI; 2003-120803/11.

Polynucleotide probes and primers for detecting beer-clouding lactic acid bacteria, for quality control during beer production applicable in bacteria, for qual brewing industry.

Claim 7; Page 31; 94pp; Japanese.

The present invention describes a polynucleotide probe, or primer, for detecting beer-clouding lactic acid bacteria containing a mucleotide sequence of (1) with 8056 base pairs (see AB276501), or a nucleotide made from not less than 15 nucleotides hybridisable with its complementary sequence. Probes and primers from the present invention can be used for detecting beer-clouding lactic acid bacteria (Lactobacillus brevis) for quality control during beer production, which is applicable in the brewing industry. The present sequence represents a PCR primer for lactobacillus brevis which is used in the exemplification of the present invention

*8888888888888

Sequence 15 BP; 4 A; 4 C; 5 G; 2 T; 0 U; 0 Other;

Gaps ô . Match 2.8%; Score 11.8; DB 1; Length 15; Local Similarity 86.7%; Pred. No. 3.8e+02; tes 13; Conservative 0; Mismatches 2; Indels 13; Conservative Query Match Matches

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RESULT 971 ABX76536/

ABX76536 standard; DNA; 15 BP.

ABX76536;

(first entry) 01-APR-2003 M. avium 23S rRNA probe #19.

Probe; 23S rRNA; 16SrRNA; tuberculosis; MTC; MOTT; peptide nucleic acid; mycobacterium tuberculosis complex; precursor rRNA; rDNA; 5S rRNA; ss; mycobacterium other than tuberculosis.

Mycobacterium avium.

US2002137035-A1.

26-SEP-2002.

07-APR-2000; 2000US-00544934.

07-APR-2000; 2000US-00544934.

(STEN/) STENDER H. (LUND/) LUND K.

(MOLL/) MOLLERUP T A.

Stender H, Lund K, Mollerup TA;

WPI; 2003-174116/17.

Peptide nucleic acid probes for detecting target sequences of Mycobacteria in samples, e.g., sputum, which are capable of hybridizing to a target sequence of mycobacterial rDNA, precursor rRNA or rRNA forming detectable hybrids.

Claim 22; Page 39; 74pp; English.

The invention relates to a peptide nucleic acid capable of hybridising to a target sequence of Mycobacterial rDNA, precursor rRNA or rRNA (55, 168 or 238) forming detectable hybrids. Also included are detecting a target sequence of mycobacteria in a sample comprising contacting rRNA or rDNA in the sample with peptide nucleic acid probes (hybridisation takes place between the probe and the rRNA or rDNA). Observing or measuring any formed detectable hybrids and relating the observation or measurement to the presence of a target sequence of mycobacteria in the sample, and a target sequence of mycobacteria in particular a target sequence of wycobacteria in particular a target sequence of complex (MrC). The probes are used for detecting a target sequence of MTC (and

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The present invention describes a method for treating, preventing or delaying neoplasm in a mammal. The method comprises administering an ErbB - 3 protein, a nucleic acid encoding an ErbB-3 protein, or their cunctional fragments, where an immune response is generated against the neoplasm. ErbB-3 has cytostatic activity, and can be used in gene theretoy. The method is useful for treating, preventing or delaying neoplasms (e.g. adrenal gland, anus, auditory nerve, bile ducts, blader, bone, brain, breast, buccal, central nervous system, cervix, colon, ear, conderving, oesophagus, eye, eyelids, fallopian tube, gastroointestinal tract, head and neck, heart, kidney, laryx, liver, lung, mandible, mandibular condyle, maxilla, mouth, nasopharym, nose, oral cavity, ovary, pancreas, parctid gland, penis, pinna, pituitary, prostate gland, covary, testes, thyroid, tonsil, uretha, uterus, vagina, ecctum, testes, thyroid, tonsil, uretha, uterus, vagina, vestibulocochlear nerve, or vulva neoplasm), or cancers (breast, ovary,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Use of an ErbB-3 protein, a nucleic acid encoding an ErbB-3 protein or their fragments, for treating, preventing or delaying neoplasms (e.g. urethra, uterus, vagina or vulva neoplasm) or cancers (e.g. breast, ovary or colon cancer).
distinguishing them from mycobacterium other than tuberculosis, MOTT) present in a sample, e.g. sputum, laryngeal swabs, gastric lavage, bronchial washings, biopsies, aspirates, expectorates, body fluids, urine, tissue sections as wall as food samples, soil, air and water samples and their cultures. The probe is able to penetrate the cell wall of the mycobacterial precursor FRNA and rRNA without harsh treatment of the mycobacterial precursor therefore avoiding a risk of interfering with the morphology of the cells. The present sequence is an M. avium probe for 16S or 23S rRNA
                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 neoplasm; ErbB-3; immune response; cytostatic; gene therapy; cancer;
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0
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                                                                                                                                                                                                                                                             Query Match 2.8%; Score 11.8; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                     Seguence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example; SEQ ID NO 9; 68pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADE36720 standard; DNA; 15 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-MAR-2003; 2003WO-CN000217.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-MAR-2002; 2002CN-00116259
                                                                                                                                                                                                                                                                                                                                                    13 AACTGCGGGTGACCG 27
                                                                                                                                                                                                                                                                                                                                                                                          15 Agcirccedericace 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-JAN-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADE36720;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADE36720
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Claim 18 claims primers for use in detecting mutations in a mammalian gene for a structural protein of cartilage comprising a sequence identified in Table I (Page 18-31). Table I includes 179 primer sequences (see AAQ65728-Q65906). The following details are given for primer 77:
Region/exon: 45 Direction: sense Primer position: 18572 (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Detecting genetic pre-disposition to osteoarthritis - and other diseases involving mutation in cartilage protein genes, by amplification and analysis of DNA and comparison with standards.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                     Gaps
stomach, prostate, colon and lung cancer). The present sequence represents an oligonucleotide used in the construction of a plasmid comprising BrbB-3, which is used in an example from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ala-Kokko L, Williams CJ, Ritvaniemi P, Baldwin C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ô
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Type II procollagen; COL2A1; amplification; primer; polymerase chain reaction; PCR; osteoarthritis; cartilage; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 2.8%; Score 11.8; DB 1; Length 16; Best Local Similarity 86.7%; Pred. No. 4.4e+02; Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                   2.8%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 3.8e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 16 BP; 4 A; 4 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                     Sequence 15 BP; 6 A; 4 C; 5 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                  Type II procollagen sequencing primer 77.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 18; Page 29; 112pp; English.
                                                                                                                                                                                                                                                                                                    AAQ65877 standard; DNA; 16 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UYJE-) UNIV JEFFERSON THOMAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  92US-00977284.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93WO-US010964.
                                                                                                                                                                                   384 GACGACGCCCAAG 398
                                                                                                                                                                                                                   1 GACGACGACGACAAG 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61 AGTCTCTGCACTACG 75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 AGTCTCTGGACTAAG 16
                                                                                                                Query Match 2.0%;
Best Local Similarity 86.7%;
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                    (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hopkingon I, Ahmad NN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1994-183530/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9411532-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-NOV-1992;
                                                                                                                                                                                                                                                                                                                                                                    25-MAR-2003
22-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-MAY-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Prockop DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                    AAQ65877;
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AAT85365
                                                                                                                                                                                                                                                                  RESULT 973
                                                                                                                                                                                                                                                                                     AAQ65877
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A novel liposomal composition of antisense oligonucleotides has been developed. The composition comprises: (a) a liposome which consists entirely of neutral phospholipids; and (b) an antisense oligonucleotide that is entrapped in the liposome and is selected from phosphodiester oligonucleotides. The present sequence represents a specifically claimed antisense p-ethoxy oligonucleotide against Sor exon 1/Abl exon 2 (B1/A2) found in human acute lymphocytic leukaemia cells. The compositions are used particularly for inhibiting the growth of tunour cells. The compositions minimise nuclease hydrolysis of the oligonucleotides and also result in increased cellular uptake and intracellular delivery of the antisense oligonucleotides. The compositions also enhance the incorporation of oligonucleotides in the liposomes compared to known liposomal formulations Anti:sense oligo:nucleotide liposomal compsns. - comprise neutral phospholipid(s) with phospho:di:ester oligo:nucleotide(s), phosphoro:thioate oligo:nucleotide(s) or p-ethoxy oligo:nucleotide(s) Gape Human; acute lymphocytic leukaemia; ALL; Philadelphia chromosome; chronic myelogenous leukaemia; Abl; break point cluster region; Bcr; inhibition; tumour; ss. ö 2.8%; Score 11.8; DB 1; Length 16; 86.7%; Pred. No. 4.4e+02; tive 0; Mismatches 2; Indels Antisense p-ethoxy oligonucleotide against leukaemia cells. Sequence 16 BP; 2 A; 4 C; 6 G; 4 T; 0 U; 0 Other; /*tag= a /note= "P-ethoxy linkages" Location/Qualifiers Claim 7; Page 19; 26pp; English. AAT85365 standard; DNA; 16 BP. AAV66874/c ID AAV66874 standard, RNA, 16 BP XX AC AAV66874; XX XX XX XX XX XX XX XX 96WO-US014146. 95US-00520385 398 GAAGGICTICIACGT 412 Lopez-Berestein G, Tari AM; 1 GAAGGCTTCTGCGT 15 (TEXA) UNIV TEXAS SYSTEM. (first entry) Local Similarity 86.7 es 13; Conservative /*tag= WPI; 1997-178904/16. 26-AUG-1996; misc_feature WO9707784-A2 29-AUG-1995; 10-DEC-1997 Synthetic AAT85365; Query Match RESULT 975 δ

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method has been developed for producing a recombinant DNA molecule. The method comprises: (a) providing a first DNA/RNA hybrid molecule comprising a first DNA linked to a first splicing component; (b) providing a second DNA/RNA hybrid molecule compriseing a second DNA/RNA hybrid molecule component is contained to that, when the first and second splicing component is selected so that, when the first and second splicing component covalently links the first and second splicing component covalently links the first DNA with the second DNA to form a dimixed together, trans-splicing to that with the second DNA to form a single, recombinant DNA molecule, and (c) admixing the first and second DNA/RNA hybrid molecules together so that the recombinant DNA molecule is produced by trans-splicing. The method can be used for the manipulation of mucleic acids. Novel genes and gene products can be generated by committed by intron sequences that can direct trans-splicing of the exon admixing nucleic acid constructs comprising exon nucleic acid sequences from a functional intron sequences the can direct trans-splicing of the exon sequences to each other. The flanking intronic sequences of the complementation between the flanking intron sequences of the complementation between the flanking intron which mediates the transesterification reactions necessary to cause the ligation of the transesterification reactions necessary to cause the ligation of the continuous nucleic acid sequences to one another, and thereby generate a recombinant gene comprised to an example from the present expenses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Manipulation of nucleic acids - using intron sequences to mediate specific cleavage and ligation of discontinuous nucleic acid molecules by trans-splicing.
                                     Human; tissue plasminogen activator; t-PA; chimeric gene assembly; manipulation; ribozyme; intron-mediated recombinant technique; cleavage; ligation; trans-splicing; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.8%; Score 11.8; DB 1; Length 16; 86.7%; Pred. No. 4.40+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tumour necrosis factor alpha antisense oligonucleotide.
Oligonucleotide for the last 16 nucleotides of K2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 16 BP; 0 A; 8 C; 3 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 36; 160pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX09083 standard; DNA; 16 BP.
                                                                                                                                                                                                                                                                                    98WO-US004881.
                                                                                                                                                                                                                                                                                                                            97US-00814412.
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nes 13, Conservative
                                                                                                                                                                                                                                                                                                                                                                      (UYBO-) UNIV BOSTON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1998-531526/45.
                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                               WO9840519-A1
                                                                                                                                                                                                                                                                                                                            11-MAR-1997;
                                                                                                                                                                                                                                            17-SEP-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                 Jarrell KA;
                                                                                                                                 Synthetic.
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Matches
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Antisense oligonuclectides (ASO) for inhibiting a tumour necrosis factoralpha (TNP-alpha) gene in an animal, preferably a human, comprise 12-50 nucleotides, 90% of which are complementary to a region of mRNA containing a GGGA sequence motif. The ASO is used to inhibit expression of a gene in an animal and for treating the animal when afflicted with a disease on disorder characterised by the presence of an mRNA from a gene containing a GGGA motif. The ASO are specifically targeted to a GGGA motif. The ASO are specifically targeted to a GGGA motif. The most efficacious ASO's contain one or more TCCC motifs. This ASO comprises a TCCC motif followed by a cytosine residue and corresponds to a region of the 1.19CAT 5' untranslated region
Tumour necrosis factor alpha; TNF-alpha; antisense oligonucleotide; ASO; inhibition; expression; treatment; disease; disorder; ss.
                                                                                                                                                                                                                                                                                                  Generation of antisense oligonucleotides - by specifically targeting a GGGA motif found in mRNA sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Recombinant nucleic acid production; combinatorial gene library; ordered gene assembly; trans-splicing; tissue plasminogen activator; kringle domain; K2; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 16 BP; 3 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kringle domain 2 (K2) nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                 Example 2; Page 37; 55pp; English.
                                                                                                                                                                                                        (UYJE-) UNIV JEFFERSON THOMAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA13286 standard; RNA; 16 BP
                                                                                                                                                                         97US-0051705P.
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99US-00101328.
                                                                                                                                            98WO-US013711.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 GATCCCCGGGTACCG 15
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                                                                                                                                                                                                                                     Tu G, Israel Y;
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                                                                              WO9901139-A1.
                                                                                                                                                                         03-JUL-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-SEP-1998;
                                                                                                             14-JAN-1999
                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA13286;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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This sequence represents a kringle domain (K2) nucleotide sequence. K2 is used in an example of the method of the invention, which demonstrates the used in an example of the method of the invention, which demonstrates the used of Englisheered ribozymes to catalyse chimeric gene assembly. The present invention relates to the in vivo production of recombinant nucleic acid sequences. The method comprises expressing in a cell, two transcripts, one contraining a first exon and first intron component, and exon. Transcript comprising a second intron components. The invention makes use of the ability of intronic sequences derived from group I or group II introns to mediate specific cleavage and ligation of discontinuous nucleic acid molecules. The method is used to produce crecombinant nucleic acid molecules. The method is used to produce into a preferably used for the preparation of exons are random. The method is preferably used for the preparation of exons are random. The method is also used for the preparation of exons are random. The method is also used for ordered and composition of exons are random. The method is also used for ordered appearation of exons are random. The method is also used for ordered and composition of exons are random. The method is also used for ordered at contacted at others. New genes can be exons but randomized at others. New genes can be exceed a specific ordered at each and a very wide range of exons may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Determining Tm range for several degenerate primers with a fixed-sequence and a degenerate-sequence portion for use in polymerase chain reaction amplification by identifying a specific sequence in the nucleic acid
                                                                                                                                                                                                                                          two RNAs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR primer; nucleic acid amplification; melting temperature; T_m; ss.
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                                                                                                                                                                                          In vivo production of nucleic acid, useful e.g. for producing combinatorial gene libraries or ribozymes, by trans-splicing t containing exon and intron component.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 2.8%; Score 11.8; DB 1; Length 16; Best Local Similarity 86.7%; Pred. No. 4.4e+02; Matches 13; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 16 BP; 0 A; 8 C; 3 G; 0 T; 5 U; 0 Other;
                                  Jarrell KA, Mulcheeva S, Donahue W;
                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 40; 186pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GENO-) GENOME TECHNOLOGIES LLC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAC63245 standard; DNA; 16 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       272 GGAGCAGGCGCCAC 286
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                                                                                                                     WPI; 2000-303208/26.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-OCT-2000.
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G; 1 T; 0 U; 0 Other;

Sequence 16 BP; 0 A; 10 C; 5

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template.
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The present invention relates to a method for selecting PCR primers for nucleic acid amplification. The method comprises determining the melting the remperature (T m) range for degenerate oligonucleotide primers with a fixed-sequence portion (FS) and a degenerate-sequence portion (DS) by searching known portion of a nucleic acid template for a sequence complementary to a desired FS of a primer. Nucleotide base pairs flanking or interspersed between the sequence complementary to a DS of one of the primers are detected and T m is calculated. The method of the present those produced. The present sequence is a primer used in the method of the present the present invention Disclosure, Fig 3A; 34pp; English.

Sequence 16 BP; 1 A; 11 C; 4 G; 0 T; 0 U; 0 Other;

Gaps . 0 2.8%; Score 11.8; DB 1; Length 16; 86.7%; Pred. No. 4.4e+02; tive 0; Mismatches 2; Indels Conservative Local Similarity 13; Query Match Matches

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142 TGGCGGTGGAGGCCG 156 15 redecedecedece g ઠે

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248/c AAC63248 standard; DNA; 16 06-FEB-2001 AAC63248; RESULT 979 AAC63248/

H

Oligonucleotide #21 used in a method for primer selection. (first entry)

PCR primer; nucleic acid amplification; melting temperature; T_m; ss.

Homo sapiens

WO200060123-A2

12-OCT-2000

05-APR-2000; 2000WO-US008962

99US-0127891P. 06-APR-1999; (GENO-) GENOME TECHNOLOGIES LLC.

Senapathy P;

WPI; 2000-656235/63

Determining Tm range for several degenerate primers with a fixed-sequence and a degenerate-sequence portion for use in polymerase chain reaction amplification by identifying a specific sequence in the nucleic acid template

Disclosure, Fig 3A; 34pp; English.

The present invention relates to a method for selecting PCR primers for mucleic acid amplification. The method comprises determining the melting temperature (T_m) range for degenerate oligonucleotide primers with a fixed-sequence portion (FS) and a degenerate sequence portion (DS) by searching known portion of a nucleic acid template for a sequence complementary to a desired FS of a primer. Nucleotide base pairs flanking or interspersed between the sequence complementary to a DS of one of the primers are detected and T m is calculated. The method of the present invention allows primers which produce more efficient DNA amplification to be produced. The present sequence is a primer used in the method of the present invention

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                                                                                                                                         therapy;
                                                                                                                                 Human; sitosterolaemia susceptibility gene; SSG; atherosclerosis; sterol-related disorder; hyperlipidaemia; hypercholesterolaemia; thera; gall stone; coronary heart disease; cardiovascular disease; arthritis; xanthoma; haemolytic anaemia; transgenic animal; ds.
                                                                                                                    Human sitosterolaemia susceptibility gene (SSG) exon5 5' splice site.
               Gaps
               ö
2.8%; Score 11.8; DB 1; Length 16; 16.7%; Pred. No. 4.4e+02; ve 0; Mismatches 2; Indels
                             305 GAGCCCCCGGGGACCG 319
        86.7%;
                                                                             AAD22030 standard; DNA; 16
                                                                                                         (first entry)
                                           15 cháccccacaccc
               Conservative
         Similarity
                                                                                                         12-FEB-2002
               13,
                                                                                            AAD22030;
   Query Match
          Local
          Best Loca
Matches
                                                                RESULT 980
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cocation/Qualifiers /*tag= a /note= "Intron" /note= "Exon" Д 10. 16 /*tag= 1 misc_feature misc_feature Key

Homo sapiens

18-APR-2001; 2001WO-US012758. WO200179272-A2 25-OCT-2001

18-APR-2000; 2000US-0198465P. 15-MAY-2000; 2000US-0204234P.

Shan B; Schultz J, Tian H,

(TULA-) TULARIK INC

WPI; 2002-017598/02.

Novel sitosterolemia susceptibility gene polypeptide and polynucleotide, useful for screening a compound that increases the level of expression o activity of SSG polypeptide for treating sterol-related disorder.

Disclosure; Fig 14B; 105pp; English.

(SSG) polypeptide. SSG is a member of adenosine triphosphate (ATP) binding cassette (ABC) family cholesterol transporter. SSG is useful for identifying a compound useful in the treatment or prevention of a strol-related disorder, including sitosterolaemia, hyperlipidaemia, astrol-related disorder, including sitosterolaemia, hyperlipidaemia, astrol-nutritichaal deficiencies. SSG is also useful for treating cholesterolaemia or conditions including coronary heart disease and other cardiovascular disease, and sitosterolaemia-associated condition including arthritis, xanthomas and chronic haemolytic anaemia. SSG expression cassette is useful in the production of transgenic non-human animals. SSG genes and their homologues are useful as tools for a number cardiovascular disorders, for forensics and paternity determinations, and for treating any of a large number of SSG associated diseases. The invention relates to an isolated Sitosterolaemia Susceptibility Gene

SXS

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The invention relates to a novel method for obtaining typing information about several variable sites within target nucleic acid, or typing one or more nucleic acid molecules. The methods of the invention are useful for typing one or more nucleic acid molecules containing two or more variable sites, preferably nucleic acid molecules containing three or more variable sites, preferably nucleic acid molecules containing three or more variable sites are typed, where three or more primer extension reactions care parformed. The method is also useful for diagnosis of pathological conditions characterized by the presence of specific nucleic acid molecule(s). The methods are particularly suited for identifying microbial species or their subtypes, and in typing procedures acid of polymorphisms, tissue typing or in clinical applications. The sequence represents a PCR primer used to sequence fragment sills of the CYP2D6 gene, which is a member of the cytochrome P450 gene superfamily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Typing nucleic acid for obtaining information about several variable sites involves simultaneously or sequentially performing two or more primer extension reactions, and determining the pattern of nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.8%; Score 11.8; DB 1; Length 16;
86.7%; Pred. No. 4.4e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                  Human, single nucleotide polymorphism; nucleic acid typing; tissue typing; sequencing; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Guanylate kinase gene associated oligonucleotide #22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 16 BP; 5 A; 3 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                               Human CYP2D6 gene sequencing primer A182FS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (PYRO-) PYROSEQUENCING AB. (STRD ) UNIV LELAND STANFORD JUNIOR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pourmand N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 5; Page 59; 86pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABX04806 standard; DNA; 16 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                            10-SEP-2001; 2001WO-GB004042.
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                                        ABN79955 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ronaghi M, Ekstroem B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-393849/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (STRD ) UNIV LELANI
(GARD/) GARDNER R.
                                                                                                                                                                                                                                                                                                                                               WO200220837-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       incorporation.
                                                                                                                                                                                                                                                                                                    Homo sapiens.
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                                                                                     ABN79955;
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ABX04806/C
ID ABX048
XC
AC ABX048
XX
DT 15-JAN
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XX
XX
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                    ABN79955
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a typing kit for judging human leukocyte antigen (HIA) genotype of a sample by hybridising a substrate on which 10-24 base oligonucleotides (AB15012-AB151809) originating in the sequences of genes e.g. belonging to HIA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HIA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when transplanting between them.
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                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; human leukocyte antigen; HLA; genotype; polymorphism; immunogenetic; transplantation; genetic disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 2.8%; Score 11.8; DB 1; Length 16; Best Local Similarity 86.7%; Pred. No. 4.4e+02; Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                            Length 16;
                                                                                       Query Match 2.8%; Score 11.8; DB 1; Length 1 Best Local Similarity 86.7%; Pred. No. 4.4e+02; Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human HLA genotyping oligonucleotide SEQ ID NO 737.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 16 BP; 2 A; 4 C; 9 G; 1 T; 0 U; 0 Other;
                                             Sequence 16 BP; 2 A; 3 C; 8 G; 3 T; 0 U; 0 Other;
present sequence is human SSG exon splice site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 10; Page 233; 345pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kagiya T, Ichihara T,
                                                                                                                                                                                                                                                                                                                                                      ABL31248 standard; DNA; 16 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JUN-2001; 2001WO-JP004662.
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                                                                                                                                                                                         142 TGGCGGTGGAGGCCG 156
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                                                                                                                                                                                                                                   1 TGCAGGTGGAGGCCG 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (NISN ) NISSHINBO IND INC. (SYST-) SYSTEM RES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
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Inoko H,

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ABL31248;

RESULT 981 ABL31248

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Gaps .. 0

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Herpesviridae; thymidine kinase; TK; DRH nucleoside binding region; viral inhibitor; bacterial inhibitor; parasite inhibitor; tumour; autoreactive immune cell; cancer; hyperkariosis; psoriasis; prostate bypertrophy; hyperthyroidism; endocrinopathy; allergy; autoimmune disease; restenosis; viral disease; AIDS; hepatitis; HCV; HBV; gene therapy; adenosine deaminase deficiency; Alzheimer's disease; guanylate kinase.
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Mus sp.

US6451571-B1

17-SEP-2002

99US-00270956 17-MAR-1999; 94US-00237592. 95US-00432871. 95US-00552304. 02-MAY-1994; 02-MAY-1995; 02-NOV-1995;

UNIW) UNIV WASHINGTON

Loeb LA, Black ME;

WPI; 2003-045581/04.

Novel Herpesviridae thymidine kinase mutant useful for inhibiting pathogens e.g. viruses, bacteria, tumor in animals, has one or more mutations encoding amino acid substitutions upstream from the DRH nucleoside binding

Example 9; Col 48; 78pp; English.

The invention describes an isolated Herpesviridae thymidine kinase (TK)
comprising a 12 amino acid (aa) nucleoside binding region having a site 3
made up of a DRH muctacoside binding site and a site 4 and mutationis, at
least one of the mutations being an aa substitution 2 or 3 aa upstream
cleast one of the mutations being an aa substitution 2 or 3 aa upstream
or for more aa downstream from the DRH motif that increases a biological
activity, preferably ability of TK to phosphorylate a nucleoside
analogue, as compared to unmutated TK. TK mutants are useful for
inhibiting pathogenic agent such as viruses, bacteria, parasites,
tumour cells or autoreactive immune cells in a warm-blooded animal. FX
mutant is useful for inhibiting a tumour or cancer in a warm-blooded
animal, for treating a variety of disease e.g., hyperkeratosis
(psoriasis), prostate hypertrophy, hyperthyroidism, endocrinopathies,
autorimmune diseases, allergies, restenosis, viral diseases such as
acquired immunodeficiency syndrome (AIDS) hepatitis (HCV or HBV),
intraccilular parasitic diseases, and to correct aberrant expression of a
cquired immunodeficiency syndrome (AIDS) hepatitis (HCV or HBV),
intraccilular parasitic diseases, and to correct aberrant expression of a
cquired immunodeficiency syndrome (AIDS) hepatitis (HCV or HBV),
intraccilular parasitic diseases, and to correct aberrant expression of a
deficiency, and Alzhaimer's diseases. The mutants are utilised as a
conditionally lethal marker for homologous recombination. This sequence
represents an oligonucleotide used in the isolation, purification and
characterisation of guanylate kinase

Sequence 16 BP; 2 A; 10 C; 2 G; 2 T; 0 U; 0 Other;

2.8%; Score 11.8; DB 1; Length 16; llarity 86.7%; Pred. No. 4.4e+02; Conservative 0; Mismatches 2; Indels Query Match Best Local Similarity Matches 13; Conserv

RESULT 984

2537/c ACD82537 standard; DNA; 16 BP

ACD82537; ACD82537/ ID ACD8: XX AC ACD8

Nucleic acid cloning associated adaptor molecule #238.

(first entry)

19-SEP-2003

nucleic acid cloning; nucleic acid ligating; mutagenesis analysis; cloning vehicle; ss. Adaptor molecule; internal deletion

Synthetic.

US2003044791-A1.

36-MAR-2003.

13-JUN-2001; 2001US-00880313

13-JUN-2001; 2001US-00880313.

(FLEM/) FLEMINGTON E

Flemington EK;

WPI; 2003-521745/49.

New adaptor molecules, useful for cloning nucleic acid molecules does not require the design and synthesis of oligonucleotides or primers.

Claim 12; Fig 5; 100pp; English.

The invention describes adaptor molecules, where each end of the adaptor is compatible with a nucleic acid digested with a restriction enzyme or a nucleic acid compatible with a nucleic acid digested with a restriction enzyme. The adaptor molecules, compositions, kits and arrays are useful for cloning nucleic acid molecules that does not require the design and synthesis of oligonucleotides or PKP primers. The adaptors, kits and arrays are also useful for ligating two ends of a single nucleic acid molecule. Or ligating two or more nucleic acid molecules. The kits can also be used for performing internal deletion mutagenesis analysis. The adaptor molecules are ligated to a cloning vehicle, making the cloning procedure more rapid and efficient, and less excreprenes. This sequence represents a nucleic acid cloning associated adaptor molecule

Sequence 16 BP; 2 A; 6 C; 5 G; 3 T; 0 U; 0 Other;

Gaps ö Match 2.8%; Score 11.8; DB 1; Length 16; Local Similarity 86.7%; Pred. No. 4.4e+02; Local 13; Conservative 0; Mismatches 2; Indels Query Match Matches

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278 GGGCGCCACCAAGCT 292 15 GGCTGCAGCAAGCT 1 ઠે 셤

AAQ47568 standard; cDNA to mRNA; 17 BP RESULT 985 AAQ47568

(revised)
(first entry) 25-MAR-2003 26-JAN-1994 AAQ47568;

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Gaps

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Specific B type jun gene probe.

quantification; human; GTP binding protein; G protein; alpha subunit; specific mRNA; detection; hybridisation; diagnosis; pathophysiology; disease state; heredditary; cancer; infectious; osteodystrophy; pituitary tumour; acromegaly; melanoma cells; diabetes; PCR; polymerase chain reaction; 88.

Synthetic

rng.res

WO9315221-A1

05-AUG-1993.

92US-00827208. 92US-00857059. 92US-00974409.

93WO-US000977

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Akitaya T, Cooper A, Mitsuhashi M;
                                  HITB ) HITACHI CHEM CO LTD.
HITB ) HITACHI CHEM RES CENT INC.
                                                   WPI; 1993-258695/32.
                    29-JAN-1992;
24-MAR-1992;
12-NOV-1992;
             29-JAN-1993;
WO9315221-A1
      05-AUG-1993
```

The sequence is that of a specific B type jun gene probe which was used in the method of the invention for the detection and quantification of manages in a sample without the need to purify the manafarom cells. The claimed method comprises identifying a polymucleotide sequence unique to the manafaron immobilising an oligomer complementary to this sequence to the minorable support the sample is then incubated with the insoluble support such that the unique sequence will hybridise to the bound cauport such that the unique sequence will hybridise to the bound of insoluble support and bound RNA is labelled in such a way that the label is incorporated onto the support relative to the amount of manafaron of used for the reliable, rapid, simultaneous quantification of multiple varieties of manafarous diseases tates, eg. hereditary diseases. Concert, and infectious diseases. G proteins are thought to be involved in causing various diseases. G proteins are thought to be involved in accountar basis of hereditary osteodystrophy. Pituleary tumours in a concentry rase also involved in invasive and metascatic melanoma cells, and dispates. See also AAQ47381-666. (Updated on 25-MAR-2003 to correct PN Quantitating messenger RNA in sample - using immobilised-polynucleotide having sequence complementary to sequence unique to the MRNA. Example 9; Page 67; 177pp; English.

Sequence 17 BP; 2 A; 6 C; 6 G; 3 T; 0 U; 0 Other;

/ Match 2.8%; Score 11.8; DB 1; Length 17; Local Similarity 86.7%; Pred. No. 5e+02; nonservative 0; Mismatches 2; Indels 0; Gaps Query Match Best Loc Matches

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8

RESULT 986 g

AAQ47593 standard; cDNA to mRNA; 17 BP. (revised)
(first entry) 25-MAR-2003 26-JAN-1994 AAQ47593; AAQ47593/C

Jun-B specific probe B-1258.

quantification; human; GTP binding protein; G protein; alpha subunit; specific mRNA; detection; hybridisation; disanosis; pathophysiology; disease state; hereditary; cancer; infectious; osteodystrophy; pituitary tumour; acromeally; melanoma cells; diabetes; PCR; polymerase chain reaction; 88.

Synthetic

The sequence is that of the jun-B specific probe B-1258 which may be used in the detection of jun oncogenes. It was used in the method of the invention for the detection and quantification of mRNAs in a sample cinvention for purify the mRNA from cells. The claimed method comprises identifying a polynuclectide sequence unique to the mRNA, and immobilises identifying a polynuclectide sequence unique to the mRNA, and comprises identifying a polynuclectide sequence unique to the mRNA, and immobilised sample is then incubated with the insoluble support such that the unique sequence will hybridise to the bound oligomer and be that the unique sequence will hybridise to the bound oligomer and be conditived. Only that the label is incorporated onto the support relative to the amount of mRNA on the support. The amount of cound label is then determined. This method can be used for the relable. Components are not be used for the relable. Components are not be used for the relative to the support of multiple varieties of mRNA. It may be used for disgnosting and recognition of pathophysiology of various diseases tates, eg. hereditation of pathophysiology of various disease states. A genetic deficiency of Gs protein is the molecular basis of hereditary osteodystrophy. Pitultary tumours in acromegalic patients have been shown component manufant Gs proteins are also involved in invasive and metastatic melanoma cells, and diabetes. See also AAQ47381-666.

Cupdated on 25-WAR-2003 to correct PN field.) Quantitating messenger RNA in sample - using immobilised-polynucleotide having sequence complementary to sequence unique to the MRNA. Gaps ; 0 Query Match

2.8%; Score 11.8; DB 1; Length 17;
Best Local Similarity 86.7%; Pred. No. 5e+02;
Matches 13; Conservative 0; Mismatches 2; Indels Sequence 17 BP; 3 A; 6 C; 6 G; 2 T; 0 U; 0 Other; Akitaya T, Cooper A, Mitsuhashi M; Example 9; Page 71; 177pp; English. (HITB) HITACHI CHEM CO LTD. (HITB) HITACHI CHEM RES CENT INC. 92US-00857208. 92US-00857059. 92US-00974409. 93WO-US000977 WPI; 1993-258695/32 29-JAN-1992; 24-MAR-1992; 12-NOV-1992; 29-JAN-1993;

217 ACTCGGTGGCGCCA 231 16 ACTTGGTGGCCGCCA 2 ò a

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(revised)
(revised)
(first entry) 14-MAY-2003 25-MAR-2003 01-SEP-1994 AAQ56954;

AAQ56954 standard; DNA; 17

RESULT 987 AAQ56954

pH 2.5 acid phosphatase probe oligo PHY-34 #3.

Phytase; pH 2.5 acid phosphatase; A. niger; strain ALK0243; mineral; liberation; phytate; plant material; feed treatment; animal; inositol; enzyme mixture; hydrolysis; phosphate; phytic acid complex; ss.

Synthetic

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The sequences given in AAOS6948-59 are probes which were used in the isolation of the pH 2.5 acid phosphatase (AP) from A. niger var. awamori strain ALKO243. These probes are based on peptide #816 and #1110 (see also ARA6255). The isolated sequences were used to transformed host calls for the expression of the pH 2.5 AP protein. The pH 2.5 AP protein in vitro, ie, in feed treatment processes, or in vivo, ie. by administering the enzymes to animals. This enzyme can be mixed to provide a balanced enzyme mixture in which comperative enzyme activity rapidly and effectively catalyses the near complete hydrolysis of phytate to incistio and free phosphate with releases of minerals from the phytic acid complex. (Updated on 25-MRA-2003 to correct PN field.) (Updated on 14-MAY-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kappa-casein; milk protein; primer; polymerase chain reaction; PCR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                            - Used to
                                                                                                                                                                                                 Turunen MK;
                                                                                                                                                                                                                                                                       Nucleic acid encoding phytase and pH 2.5 acid phosphatase - Used to produce the enzymes and enzyme mixts. for liberating minerals from phytate, partic. for animal feed.
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.8%; Score 11.8; DB 1; Length 17; 86.7%; Pred, No. 5e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                 Paloheimo MT, Fagerstroem RB, Miettinen-Oinonen ASK, Rambosek JA, Piddington CS, Houston CS, Cantrell MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 5 A; 6 C; 0 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                       Example 1; Page 31; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kappa-casein DNA primer AA75-80B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (COLS ) UNIV COLORADO FOUND INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ89601 standard; DNA; 17 BP
                                                                          93WO-US007058.
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                                                                                                        92US-00925401.
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                                                                                                                                                 NEVALAINEN H K M.
                                                                                                                                                                                                                                              WPI; 1994-065302/08.
                                                                                                                                      PANL-) PANLABS INC
                                                                                                                                                                     ALKO LTD
                WO9403072-A1
                                                                        27-JUL-1993;
                                                                                                        31-JUL-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3-0CT-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-OCT-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-NOV-1995
                                            17-FEB-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US5391497-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-FEB-1995,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ89601;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                    (NEVA/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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Enzymatic nucleic acid, ribozyme, trans cleavage; inhibition, devargedulation, interleukin-5; IL-5; ICAM-1; intercellular adhesion molecule; rel A; tumour necrosis factor; riberalpha, respiratory syncytial virus, RSV; bcr-abl; oncogene; rranslocation; chronic myelogenous leukaemia; CML; cancer; Philadelphia chromosome; inflammation; autoimmune disease; atherosclerosis myocardial infarction; stroke; restenosis; atherosclerosis; myocardial infarction; stroke; restenosis; myocardial infarction; stroke; restenosis; myocardial; Kawasaki disease; septic shock; HIV; human immunodeficiency virus; acquired immune deficiency syndrome; AlDS;
                                                                                                                                                         A commercial cDNA library prepd. in lambda gt11 from mRNA obtd. from human breast tissue removed during the third trimester of pregnancy was screened with rabbit anti-bovine kappa-casein cDNA. The cDNA insert of a recombinant phage was amplified by PCR using the lambda sequencing primers given in AAQ89599-600 and the kappa-casein primer given in AAQ8959-600 and the kappa-casein primer given in AAQ89501 to obtain a clone encoding human kappa-casein
                                                                            DNA encoding human kappa-casein - used for the prodn. of large amts. of highly purified kappa-casein milk protein for infant use.
                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rat ICAM hammerhead ribozyme target sequence (nt. position 1092).
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                                                                                                                                                                                                                                                                                                       Query Match
2.8%; Score 11.8; DB 1; Length 17;
Best Local Similarity 86.7%; Pred. No. 5e+02;
Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                            Sequence 17 BP; 4 A; 3 C; 7 G; 3 T; 0 U; 0 Other;
Chang Y;
                                                                                                                               Example D; Col 11; 14pp; English.
 Menon RS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT53541 standard; RNA; 17 BP.
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94US-00218934.
94US-00222795.
94US-00224483.
94US-00224958.
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94US-00271280.
94US-00291932.
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94US-00292620.
94US-00293520.
94US-00300000.
94US-00303039.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (revised)
(first entry)
 Jeffers KF,
                               WPI; 1995-160470/21.
                                                 P-PSDB; AAR72699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rattus rattus.
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27-MAR-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-APR-1994
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 Ham RG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 989
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Rattus rattus.

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The present sequence represents a preferred target sequence for an enzymatic nucleic acid (i.e. a ribozyme) which cleaves ICAM-1 mRNA at the nucleotide base position indicated in the DE line. Regions of the mRNA that do not form secondary folding structures and that contain potential hammerhead and hairfin ribozyme cleavage sites were identified by computer analysis. Ribozymes directed against these mRNA sequences were designed and synthesised with modifications that improve their nuclease resistance. The ribozymes cleave the ICAM-1 target sequences and thereby inhibit ICAM-1 expression, making them useful for reducing transplant rejection and alleviating symptoms in patients with rheumatoid arthritis, asthma and other inflammatory disorders. (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Enzymatic nucleic acid; ribozyme; trans cleavage; inhibition; gene expression; downregulation; interleukin-5; IL-5; ICAM-1; intercellular adhesion molecule; rel A; tumour necrosis factor; TNF-alpha; respiratory syncytial virus; RSV; bcr-abl; oncogene; transbocation; chronic myelogenous leuksemia; (ML; cancer; Philadelphia chromosome; inflammation; autoimmume disease; atherosclerosis; myocardial infarction; stroke; restenosis; transplant rejection; heumatoid arthrits; psorifasis; myocardial ischaemia; Kawasaki disease; septic shock; HIV; human immunodeficiency virus; acquired immune deficiency syndrome; AIDS;
                                                                                                                                                                                                                                                                                  DDT, Chowrira B, Direnzo A, Draper KG, Dudycz LW;
Karpeisky A, Kisich K, Matulic-Adamic J, Mcswiggen JA;
Pavco P, Beigleman L, Sullivan SW, Sweedler D, Thompson JD;
Usman N, Wincott FE, Woolf T;
                                                                                                                                                                                                                                                                                                                                                                                                                    Ribozymes having modified bases and methods for producing them - for use in inhibiting disease related genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    / Match 2.8%; Score 11.8; DB 1; Length 17; Local Similarity 73.3%; Pred. No. Se+02; nes 11; Conservative 2; Mismatches 2; Indels 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rat ICAM hammerhead ribozyme target sequence (nt. position 1295).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 5 A; 3 C; 7 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 2; Page 202; 407pp; English.
                94US-00311749
94US-00314377-
94US-00319492-
94US-00319493-
94US-00334847-
94US-0033516-
94US-00345516-
94US-00345516-
94US-00345518-
94US-00345518-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 AGAGGGUCUCAGCA 15
                                                                                                                                                                                                                                                 (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-MAR-2003 (revised)
27-MAR-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                                                               WPI; 1995-351090/45.
                                                                                                                                                                                                                                                                                    Stinchcomb DT,
                                                     03-OCT-1994;
07-OCT-1994;
11-OCT-1994;
                                                                                                                                                                                     23-DEC-1994;
                                                                                                               04-NOV-1994;
                                                                                                                                                      28-NOV-1994;
                                                                                                                                                                       16-DEC-1994;
                    23-SEP-1994
                                                                                                                                    10-NOV-1994
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Modak A,
Tracz D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT53575
244454444
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The present sequence represents a preferred target sequence for an enzymatic nucleic acid (i.e. a ribozyme) which cleaves ICAM-1 mRNA at the nucleotide base position indicated in the DE line. Regions of the mRNA that do not form secondary folding structures and that contain potential hammerhead and hairpin ribozyme cleavage sites were identified by computer analysis. Ribozymes directed against these mRNA sequences were designed and synthesised with modifications that improve their nuclease resistence. The ribozymes cleave the ICAM-1 target sequences and thereby inhibit ICAM-1 expression, making them useful for reducing transplant rejection and alleviating symptoms in patients with rheumatoid arthritis, asthma and other inflammatory disorders. (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      b DT, Chowrira B, Direnzo A, Draper KG, Dudycz LW; Karpelsky A, Klaitch K, Matulic-Adamic J, Mcswiggen JA; Pavco P, Beigleman L, Sullivan SM, Sweedler D, Thompson JD; Usman N, Wincott FE, Woolf T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ribozymes having modified bases and methods for producing them - for use in inhibiting disease related genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.8%; Score 11.8; DB 1; Length 17; 73.3%; Pred. No. 5e+02; Live 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 5 A; 3 C; 7 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 2; Page 202; 407pp; English.
                                                                                                                                 9405 00218934
9408 00222438
9408 00222483
9408 00228641
9408 00271280
9408 00271280
9408 00291633
9408 00291630
9408 00391680
9408 0031486
9408 0031486
9408 0031486
9408 0031489
9408 0031489
9408 0031489
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                                                                                            95WO-IB000156
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 73.3
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1995-351090/45.
                                        WO9523225-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-JAN-1995;
                                                                                            23-FEB-1995;
                                                                   31-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Stinchcomb
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Modak A,
Tracz D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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à
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W09531571-A2.

02-JUL-1996

AAT06812;

RESULT 991 AAT06812 04-MAY-1995;

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A set of internal sequencing primers (AAT35281-91) were used to sequence CDNA clone E1-C19 (see also AAT35277), which codes for chemokine receptor K5.5 (AAR99274). They were designed on the basis of previous sequencing results
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chemokine receptor which binds MIP-1-alpha, RANTES and/or MCP-1 - useful in screening for agents to treat asthma, hay fever, eczema, allergies, atopic dermatitis, rhinitis or conjunctivits.
                                                                                                   Chemokine receptor K5.5; MIP-1-alpha, RANTES; MCP-1; allergy; atheroma; HIV; AIDS; graft rejection; stem cell; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Probe A' (Set 9) for M. tuberculosis 16S rRNA gene nucleotides 721-760.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                probe, modified ligase chain reaction, Mycobacterium tuberculosis, M. avium; M. intracellulare, M. kansasii; detection; diagnosis; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Semplefacey IE, Manlove MT, Solomon NA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.8%; Score 11.8; DB 1; Length 17;
86.7%; Pred. No. 5e+02;
iive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 3 A; 5 C; 5 G; 4 T; 0 U; 0 Other;
                                                            Chemokine receptor K5.5 primer K5-5D (sense).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example; Fig 2; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT06537 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                              96WO-GB000143.
                                                                                                                                                                                                                                                                                                                                                       95GB-00001683.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  94US-00242403
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(first entry)
                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.8%
Query Match
Best Local Similarity 86.7%
Matches 13, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                (GLAX ) GLAXO GROUP LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Leckie GW, Davis AH,
                                                                                                                                                                                                                                                                                                                                                                                                                                              Wells TNC, Power CA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1996-362692/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ABBO ) ABBOTT LAB.
                                                                                                                                                                                                                                                                                                              24-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-MAY-1995;
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                                                                                                                                                                                                                       WO9623068-A1.
                                                                                                                                                                                                                                                                                                                                                       27-JAN-1995;
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02-JUL-1996
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                   09-DEC-1996
                                                                                                                                                                                                                                                                  01-AUG-1996.
                                                                                                                                                                           Synthetic.
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셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                probe set 9 (AATO6811-814) were selected to detect a target sequence in the 16S ribosomal RNA gene (nucleotides 721-760) of M. uberculosis. The detecting target DNA from several species of bacteria of the genue (mucleotides 721-760) of M. uberculosis. The detecting target DNA from several species of bacteria of the genue (Mycobeacterium. A modified ligase chain reaction was utilised which uses two pairs of probes designated A. B. (primary probes) and A., is (secondary probes). Probe pairs were directed to the same target strand and ultimately ligated to one another after annealing to the target strand and ultimately ligated to one another after annealing to the target create a gap between one probe terminus and then east probe terminus when the pair was annealed to the target sequence. Other modified ends include a base mismatched with the target sequence. Other modified ends include a base mismatched with the target sequence. The presence of modified ends a base mismatched with the signal oreated by blunt-end ligation of the complementary probe duplaces to one another in the absence of target. Correction of the modification, in a target dependent manner, was the fused (reorganised) probe was dissociated (e.g. melted) from the target and, as with conventional LCR, the process was repeated for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New probes for detection of Mycobacterium species - derived from the 16S ribosomal RNA gene, the protein antigen b gene and the 65 kD and 10 kD heat shock protein genes of M.tuberculosis.
                                                                                                                                                                                              Probe A' (Set 9) for M. tuberculosis 16S rRNA gene nucleotides 721-760.
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                                                                                                                                                                                                                                          probe; modified ligase chain reaction; Mycobacterium tuberculosis; M. avium; M. intracellulare; M. kansasii; detection; diagnosis; ss.
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86.7%; Pred. No. 5e+02;
rative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kratochvil JD, Leckie GW, Odonnell DL, Solomon NA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 2 A; 9 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 2; Page 41; 60pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAT35286 standard; DNA; 17 BP.
                                                            AAT06812 standard; DNA; 17 BP.
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                                                                                                                                                                                                                                                                                                              Mycobacterium tuberculosis.
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AAT35286;

RESULT 992 AAT35286/c ID AAT352 XX AC AAT3521

Query Match

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Gaps

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New probes for detection of M.tuberculosis - derived from e.g. the gene coding for protein antigen b and from the insertion-like element 186110 of M.tuberculosis.

Example 8; Page 41; 60pp; English.

probe set 9 (AAT06536-539) were selected to detect a target sequence in the 16S ribosomal RNA gene (nucleotides 721-760) of M. uberculosis. The probes were labelled with blotin and fluorescain. Set 9 as capable of detecting target DNA from several species of bacteria of the genus Mycobacterium. A modified ligase chain reaction was utilised which uses two pairs of probes of primary probes) and A., B. (primary probes) and A. B. (secondary probes). Probe pairs were directed to the same target strand and ultimately ligated to one another after annealing to the target strand and ultimately ligated to one another after annealing to the target service of a pair and and bases omitted to respect to the point of 14gation. The modified end had bases omitted to create a gap between one probe terminus and the next probe terminus when the pair was annealed to the target sequence. Other modified ends include a base mismatched with the target sequence. Other modified ends include a base mismatched with the target sequence. The presence of modified ends include complementary probe duplexes to one another in the absence of target. Corpiementary probe duplexes to one another in the absence of target. Corpiementary probe duplexes to one another in the absence of target. The fused (fresongaised probe was dissociated (e.g. melted) from the target and, as with conventional LCR, the process was repeated for target and, as with conventional LCR, the process was repeated for

Sequence 17 BP; 2 A; 9 C; 4 G; 2 T; 0 U; 0 Other;

0; Gaps 2.8%; Score 11.8; DB 1; Length 17; 86.7%; Pred. No. 5e+02; ative 0; Mismatches 2; Indels Query Match Best Local Similarity 86.7¹ Matches 13, Conservative

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330 GCGGACGACCAGGGC 344

AAT62817 standard; DNA; 17 BP. AAT62817; AAT62817

(first entry) 18-NOV-1997

Primer MGHR1 for murine growth hormone cDNA.

Primer; polymerase chain reaction; PCR; amplification; murine; mouse; growth hormone; transformation; stem cell; mammal; transformed organism; increased growth; continuous expression; improvement; body weight; milk production; ss.

Synthetic.

WO9708947-A1

13-MAR-1997.

96WO-JP002402. 28-AUG-1996; 95JP-00231086. 08-SEP-1995;

(TAKI) TAKARA SHUZO CO LTD.

Kato I; Matsushita H, Asada K, Okado T, Zhang Y,

WPI; 1997-192587/17.

Organisms transformed by growth hormone gene - for producing higher body

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The present sequence is a primer for the PCR amplification of murine growth hormone (mGH) cDNA, which was used to transform a stem cell, which in turn was introduced into an organism to produce a transformed organism. The transformed organism exhibits increased growth, and as the growth hormone gene is expressed continuously, it can be grown very quickly. The resulting organism, specifically a mammal, shows improved
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                                                                                                                                                                                                                                                                             Sequence 17 BP; 4 A; 3 C; 10 G; 0 T; 0 U; 0 Other;
                                            Example 1; Page 22; 39pp; Japanese.
weight, faster growing specimens.
                                                                                                                                                                                                                                body weight and milk production
                                                                                                                                                                                                                                                                                                                                                                                                                   272 GGAGCAGGGCGCAC 286
                                                                                                                                                                                                                                                                                                                         Query Match
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Gaps

AAX68713/c ID AAX68713 standard; RNA; 17 BP. (first entry) 28-JUL-1999 AAX68713; RESULT 995

1 GGGGCAGGGAGGCAC 15

Human flt1 VEGF receptor hammerhead ribozyme substrate #8.

Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead riboxyme; hairpin riboxyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinsse 1; kinase insert domain containing receptor; foetal liver kinase 1; ss.

Homo sapiens.

WO9715662-A2.

96WO-US017480. 25-OCT-1996; 01-MAY-1997.

95US-0005974P. 96US-00584040. 26-OCT-1995; 11-JAN-1996;

RIBO-) RIBOZYME PHARM INC. CHIR) CHIRON CORP

Mcswiggen J, Stinchcomb D, Escobedo J; Pavco P,

WPI; 1997-259017/23.

Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.

Claim 4; Page 46; 218pp; English.

The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VBGF). A patient (preferably human) having a condition associated with the level of the fina-like tyrosine kinase 1 (fit-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (fik-1) (e.g. tumour angiogenesis, ocular disease, psoriasis and rheumatoid arthritis) can be tracted by administering the nucleic acid molecule or the expression vector to the patient. AXX67275 to AXX7752 represent specific examples of nucleic acid molecules from the present invention

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Wed Apr 21 12:58:21 2004
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AAX68723 standard; RNA; 17 BP.

RESULT 99 AAX68723/ ID AAX6

aa i Gui

(first entry)

28-JUL-1999

AAX68723;

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The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fims-like tyrosine kinase 1 (fll-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. turmour anglogenesis, contlar diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the parient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention
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                                                                                                                                                                                                                                                                                                                                                               Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour anglogenessis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
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86.7%; Pred. No. 5e+02;
iive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                    Human flt1 VEGF receptor hammerhead ribozyme substrate #541.
                                    / Match 2.8%; Score 11.8; DB 1; Length 17; Local Similarity 86.7%; Pred. No. 5e+02; les 13; Conservative 0; Mismatches 2; Indels
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          BP; 1 A; 5 C; 9 G; 0 T; 2 U; 0 Other;
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96US-00584040.
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                                                                                                                 299 GGACCTGAGCCCCGG 313
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                                                                                                                                                15 gcacccaacccccc 1
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11-JAN-1996;
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              Sequence 17
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The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (WGRF). A parient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (fll-1), kinase insert domain containing receptor (XDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, coular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
                                                                                                     Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; templike; tyrosine kinase 1; kinase insert domain containing receptor;
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                                                                         Human flt1 VEGF receptor hammerhead ribozyme substrate #18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mouse flt-1 VEGF receptor hammerhead ribozyme substrate
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96US-00584040.
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Best Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                                             fms-like tyrosine kinase
foetal liver kinase 1; ss
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11-JAN-1996;
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247 TCCCGGGCTCGGCCA 261

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13;

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Similarity

17 rccceeccaaecca 3

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The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (WEGF). A patient (preferably human) having a condition associated with the level of the fims-like tyrosine kines 1 (fll-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour anglogenesis, coular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
Vascular endothelial growth factor receptor; VBGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour anglogenessis; psoriasis; rheumatoid arthritis; ocular disease; fms-1ike tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 / Match 2.8%; Score 11.8; DB 1; Length 17; Local Similarity 86.7%; Pred. No. 5e+02; les 13; Conservative 0; Mismatches 2; Indels 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                          Stinchcomb D, Escobedo J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 1 A; 7 C; 5 G; 0 T; 4 U; 0 Other;
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96US-00584040.
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(CHIR ) CHIRON CORP.
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11-JAN-1996;
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The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VGZF). A patient (preferably human having a condition associated with the level of the fms-like tyrosine kinase 1 (Flt-1), kinase insert domain containing receptor (XDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, coular diseases, postriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention

Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.

Claim 4; Page 56; 218pp; English

Escopedo J;

Stinchcomb D,

Pavco P, Mcswiggen J, WPI; 1997-259017/23

95US-0005974P. 96US-00584040. 96WO-US017480,

11-JAN-1996; 25-OCT-1996; 26-OCT-1995;

(RIBO-) RIBOZYME PHARM INC. (CHIR) CHIRON CORP.

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                                                                                                                                                                                                                                                                                                                                                                                                                                       Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; rumour anglogenessis; psoriasis; rheumatoid arthritis; ocular disease; fms-1ike tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; ss.
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96US-00584040.
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                                                                                                                                                                                                                                                                                                             43 ATGGCCACCACTCAG 57
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Matches 11; Conservative
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11-JAN-1996;
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Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor;

Foetal liver kinase 1; ss

Homo sapiens WO9715662-A2

01-MAY-1997

Human flt1 VEGF receptor hammerhead ribozyme substrate #339.

28-JUL-1999 (first entry)

AAX69044;

AAX69044 standard; RNA; 17 BP

RESULT 999

AAX69044

WPI; 1997-259017/23

Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.

Claim 4; Page 155; 218pp; English.

The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VGGF). A patient (preferably human) having a condition associated with the level of the fine like tyrosine kinase 1 (fit.1), kinase insert domain containing receptor (XDR) and/or fortal liver kinase 1 (fik.1) (e.g. tumour receptor (XDR) and/or fortal liver kinase 1 (fik.1) (e.g. tumour receptor (XDR) by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention

Sequence 17 BP; 2 A; 8 C; 3 G; 0 T; 4 U; 0 Other;

0; Gaps 2.8%; Score 11.8; DB 1; Length 17; 86.7%; Pred. No. 5e+02; tive 0; Mismatches 2; Indels Best Local Similarity 86.73 Matches 13; Conservative Query Match

411 GIGATCGAGACGCGG 425 15 Greadcaacaccec 1 a

AAT76176 standard; DNA; 17 BP RESULT 1001 AAT76176 ID AAT76176

(first entry) 12-SEP-1997

AAT76176;

Human IL3 receptor antisense oligonucleotide.

Asthma; airway epithelium; adenosine free; cystic fibrosis; chronic obstructive pulmonary disease; bronchitis; interleukin; ss.

WO9640162-A1

19-DEC-1996.

96WO-US009306. 06-JUN-1996; 95US-00474497. 07-JUN-1995;

(UYEC-) UNIV EAST CAROLINA.

Metzger WJ; Nyce JW,

WPI; 1997-051871/05.

Treatment of airway diseases such as asthma - by topically applying adenosine-free antisense oligo:nucleotide to airway epithelium of subject.

Example 5; Page 29; 71pp; English.

A method for treating airway disease in a subject has been produced, which involves the topical administration of an essentially adenosine free antisense oligonucleotide (ON) to the airway epithelium of the subject. The present sequence is an antisense oligonucleotide specific for the human IL3 receptor. The method can be used to treat airway diseases such as cystic fibrosis, asthma, chronic obstructive pulmonary disease, bronchitis and other airway diseases characterised by an inflammatory response. By eliminating adenosine from the antisense ON,

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its liberation upon antisense degradation is prevented, thereby preventing adenosine-induced bronchoconstriction in patients with hyper-reactive airways
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                                                                                                        2.8%; Score 11.8; DB 1; Length 17;
86.7%; Pred. No. 5e+02;
tive 0; Mismatches 2; Indels
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RESULT 1002

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AAA22825 standard; RNA; 17 BP AAA22825

AAA22825;

19-JUN-2000 (first entry)

Integrin subunit beta 3 substrate sequence SEQ ID NO:6051.

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Human; aryl hydrocarbon nuclear transport; ARNY; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; antidiabetic; ophthalmologic; antiliflammatory; antiarchritic; antidiabetic; dermatological; RNA cleavage; cencer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; verruea vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; ss.

Homo sapiens.

07-0CT-1999.

99WO-US006507. 24-MAR-1999;

(RIBO-) RIBOZYME PHARM INC.

Pavco PA, Roberts E, Jarvis T, Coeshott C, Mcswiggen JA;

WPI; 1999-591315/50.

Novel ribozymes for modulating the synthesis, expression and/or stability of an mRNA encoding an angiogenic factors.

Claim 54; Page 244; 305pp; English.

The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an arryl hydrocarbon nuclear transporter (ARNI) gene, an integrin subunit beta 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA1675 to AAA1167 and AAA1761 to AAA1762 represent ribozyme sequences for ARNIT, and AAA1166 to AAA1762 represent ribozyme sequences for ARNIT, and AAA19154 represent their corresponding target sequences, for Tie-2, and AAA19167 to AAA19168 to AAA19169 represent ribozyme sequences; AAA19169 to AAA2169 and AAA2169 represent ribozyme sequences; AAA19169 to AAA2169 and AAA2169 to AAA2342 to AAA2343 to AAA23342 represent ribozyme sequence; for integrin subunit beta 3, and AAA2346 to AAA23343 to Present ribozyme sequence; the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding anglogenic factor, especially ARNI,

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integrin subunit beta-3, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related macular degeneration (ARMD), inflammation, and arthritis, as well as neovascular glaucoma, myopic degeneration, psoriasis, verruca vulgaris, angiofibroma of tubercus sclerosis, pot wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, and other syndromes and diseases related to the levels of ARMT, Tie-2, integrin subunit alpha-6, or integrin subunit beta-3
         88888888888888
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Sequence 17 BP; 0 A; 8 C; 4 G; 0 T; 5 U; 0 Other;

2.8%; Score 11.8; DB 1; Length 17; 66.7%; Pred. No. 5e+02; ative 3; Mismatches 2; Indels 0; Gaps 242 CTGCTTCCCGGGCTC 256 3 ccecurcccgeguuc 17 ων.ν. Best Local Similarity 66.7# Matches 10; Conservative ठे

AAA22832 standard, RNA; 17 BP. 19-JUN-2000 (first entry) AAA22832; RESULT 1003 AAA22832,

Integrin subunit beta 3 substrate sequence SEQ ID NO:6058.

Human, aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antidiabetic; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tubercous sclerosis; pot-wine stain; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; ss.

Homo sapiens

07-OCT-1999.

99WO-US006507. 24-MAR-1999;

(RIBO-) RIBOZYME PHARM INC.

Pavco PA, Roberts E, Jarvis T, Coeshott C, Mcswiggen JA;

WPI; 1999-591315/50.

Novel ribozymes for modulating the synthesis, expression and/or stability of an mRNA encoding an anglogenic factors. Claim 54; Page 245; 305pp; English.

The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNY) gene, an integrin subunit beta 3 gene, an integrin subunit beta 3 gene, an integrin subunit beta 3 gene, an AAA1168 to AAA1762 represent ribozyme sequences for AAA16775 to AAA1168 to AAA1765 to AAA1762 represent ribozyme sequences for AAA19087 to AAA19154 represent ribozyme sequences, AAA1685 to AAA18386 and AAA19087 to AAA19155 to AAA1922 represent their corresponding target sequences for Tie-2, and AAA19386 to AAA19086 and AAA19155 to AAA1925 represent their corresponding target sequences; sand AAA1925 to AAA1925 in their ribozyme sequences for integrin alpha 6 subunit, and AAA2362 to AAA21500 and AAA21596 to AAA21698 represent their corresponding target sequences;

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AAA21669 to AAA22475 and AAA23263 to AAA23342 represent ribozyme sequence for integrin subunit beta 3, and AAA22476 to AAA23262, AAA23343 to AAA23422 represent their corresponding target sequences. The ribozymes of the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially ARNT, integrin subunit beta-3, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related necovaccular glaucoma, myopic degeneration, and arthritis, as well as necovaccular glaucoma, myopic degeneration, psoriasis, verruca vulgaris, and ofther syndrome, and arthritis as well as and other syndrome, and arthritis as well as and other syndrome, and etales of ARNT, Tie-2, integrin subunit beta-3
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Sequence 17 BP; 3 A; 7 C; 4 G; 0 T; 3 U; 0 Other;

0; Gaps Query Match 2.8%; Score 11.8; DB 1; Length 17; Best Local Similarity 86.7%; Pred. No. 5e+02; Matches 13; Conservative 0; Mismatches 2; Indels

230 CAAATCGGGAGGCTG 244 17 cracressesers 3 a

AAA21483 standard; RNA; 17 BP. RESULT 1004 AAA21483,

Integrin alpha 6 subunit substrate sequence SEQ ID NO:4709.

19-JUN-2000 (first entry)

AAA21483;

Human, aryl hydrocarbon nuclear transport; ARNT, TIB-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme, angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARMD; dermacological; MNA cleavage, cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tubercous sclerosis; pot-wine stath; Sturge Waber syndrome; Kippel-Trenaunay-Waber syndrome; ss.

Homo sapiens.

WO9950403-A2.

07-OCT-1999.

99WO-US006507. 24-MAR-1999;

98US-0079678P. 27-MAR-1998;

(RIBO-) RIBOZYME PHARM INC.

Pavco PA, Roberts E, Jarvis T, Coeshott C, Mcswiggen JA;

WPI; 1999-591315/50.

Novel ribozymes for modulating the synthesis, expression and/or stability of an mRNA encoding an angiogenic factors.

Claim 55; Page 211; 305pp; English.

The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an arryl hydrocarbon nuclear transporter (ARNY) gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA1675 to AAA17661 to AAA1767 represent ribozyme sequences for ARNT, and AAA17661 and AAA17662 to AAA17684 represent their corresponding target sequences; AAA17684 represent their corresponding target sequences; AAA17684 represent thouse

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AAA19154 represent ribozyme sequences for Tie-2, and AAA18386 to AAA19086
and AAA19155 to AAA21822 represent their corresponding target sequences;
AAA19125 to AAA21861 and AAA21801 to AAA21805 represent ribozyme
sequences for integrin alpha 6 subunit, and AAA21805 to AAA21500 and
AAA2189 to AAA2186 and AAA22325 to AAA21362 represent ribozyme sequence;
for integrin subunit beta 3, and AAA22476 to AAA21362, AAA21343 to
AAA21422 represent their corresponding target sequences. The ribozymes of
the invention are used for modulating the synthesis, expression and/or
stability of an mRNA encoding angiogenic factor, especially ARNT,
integrin subunit beta-3, integrin subunit alpha-6, or Tie-2. They are
especially used to treat cancer, diabetic retinopathy, age related
macular degeneration (ARND), inflammation, and arthritis, as well as
neovascular glaucoma, myopic degeneration, psoriasis, verruca vulgaris,
angiofibroma of tuberous sclerosis, pot-vine stains, sturge Weber
syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome,
and other syndromes and diseases related to the levels of ARNT, Tie-2,
integrin subunit alpha-6, or integrin subunit beta-3
             88888888888888888888888888888
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Sequence 17 BP; 6 A; 1 C; 4 G; 0 T; 6 U; 0 Other;

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0; Gaps
Match 2.8%; Score 11.8; DB 1; Length 17; Local Similarity 86.7%; Pred. No. Se+02; Les 13; Conservative 0; Mismatches 2; Indels
        Query Match
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AAA22734 standard; RNA; 17 BP. AAA22734; RESULT 1005

19-JUN-2000 (first entry)

Integrin subunit beta 3 substrate sequence SEQ ID NO:5960.

Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; harmenead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; vertuce vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; ss.

Homo sapiens.

WO9950403-A2.

99WO-US006507. 24-MAR-1999;

98US-0079678P. 27-MAR-1998;

(RIBO-) RIBOZYME PHARM INC.

Pavco PA, Roberts E, Jarvis T, Coeshott C, Mcswiggen JA;

98WO-US019419. 97US-0059160P 98US-00093972.

17-SEP-1998; 17-SEP-1997; 09-JUN-1998; (UYEC-) UNIV RAST CAROLINA.

WPI; 1999-229400/19.

Nyce JW;

WPI; 1999-591315/50.

Novel ribozymes for modulating the synthesis, expression and/or stability of an mRNA encoding an anglogenic factors.

Claim 54; Page 239; 305pp; English.

The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl

hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene, AAA16775 to AAA1761 to AAA1762 represent ribozyme sequences for ARNT, and AAA17168 to AAA17623 to AAA1889 represent their corresponding target sequences; AAA17623 to AAA18885 and AAA18987 to AAA19154 represent ribozyme sequences for Tie-2, and AAA18968 to AAA19155 to AAA19155 to AAA19168 to AAA19168 to AAA19168 to AAA19168 to AAA19168 to AAA2186 and AAA19168 to AAA2186 and AAA2180 to AAA2180 and AAA2180 to ô Antisense oligonucleotide; multiple target; antisense treatment; impaired respiration; inflammation; lung disease; pulmonary vasconstriction; inflammation; allergic rhinitis; acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; pulmonary hypertension; pulmonary vasconstriction; emphysema; cohronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pencreatic cancer; hepatocellular carcinoma; kidney cancer; melanoma; hepatic metastasis; 2; Indels 0; Gaps Query Match 2.8%; Score 11.8; DB 1; Length 17; Best Local Similarity 86.7%; Pred. No. 5e+02; Matches 13; Conservative 0; Mismatches 2; Indels Human IL-3 receptor antisense oligonucleotide fragment. integrin subunit alpha-6, or integrin subunit beta-3 Sequence 17 BP; 3 A; 7 C; 4 G; 0 T; 3 U; 0 Other; AAX53973 standard; DNA; 17 BP. 230 CAAATCGGGAGGCTG 244 17 CTACTCGGGAGGCTG 3 05-JUL-1999 (first entry) prostate cancer; ss. WO9913886-A1 Synthetic. AAX53973; RESULT 1006 AAX53973 ઠ

Wed Apr 21 12:58:21 2004

Disclosure; Page 31; 75pp; English.

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The specification describes antisense oligonucleotides (AAX52869-X55271)

directed against at least 2 mRNAs selected from target genes, coding and
non-coding regions of RNAs corresponding to target genes, genes, interation
codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'
end and the juxta-section between coding and non-coding regions and all
segments of RNAs encoding proteins associated with one or more diseases,
conditions or mixtures. The antisense oligonucleotides may be derived
from sequences AAX55272-74. These multiple target oligonucleotides
conditions or mixtures. The antisense oligonucleotides may be derived
from sequences AAX55272-74. These multiple target oligonucleotides
conditions. Typical diseases and conditions are those
associated with impaired respiration and inflammation, including lung
diseases, pulmonary vasconomeriation, inflammation, relation relation,
acute asthma, allergies, asthma, impeded respiration, respiratory
distress syndrome, pain, cystic fibrosis, pulmonary hypertension,
distress syndrome, pain, cystic fibrosis, pulmonary hypertension,
disease (COPD), and cancers such as leukemias, lymphomas, carcinomas e-g
colon cancer, breast cancer, lung cancer, pancreatic cancer,
colon cancer, breast cancer, lung cancer, pancreatic cancer,
colon cancer, lung cancer, metastasize or have metastasized
colon target, including breast and prostate cancer New antisense oligonucleotides used in treatment of, e.g. pulmonary vasoconstriction. Disclosure; Page 48; 120pp; English.

Sequence 17 BP; 0 A; 7 C; 3 G; 7 T; 0 U; 0 Other;

0; Gaps 2.8%; Score 11.8; DB 1; Length 17; 86.7%; Pred. No. 5e+02; ive 0; Mismatches 2; Indels 13; Conservative Local Similarity Matches

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242 CIGCIICCCGGGCIC 256 CICITICCCGGGCIC 15 ઠે a

AAV72257 standard; DNA; 17 BP RESULT 1007 AAV72257,

24-MAY-1999 (first entry) AAV72257;

Gal4 binding site; gene expression regulation; transgenic plant; chimeric; promoter; DNA binding domain; plant disease resistance; pest resistance; grain quality; oil composition; starch composition; protein composition, transcription factor; seed storage protein; multiple transgene regulation; tissue-specific promoter; developmentally regulated promoter; so. cerevisiae galactose metabolism gene promoter Gal4 binding site.

Saccharomyces cerevisiae.

WO9859062-A1.

30-DEC-1998

98WO-US013006. 23-JUN-1998; 97US-00881687 24-JUN-1997; (DUPO) DU PONT DE NEMOURS & CO E I.

Odell JT; Liu Z,

WPI; 1999-105629/09.

Regulating gene expression in a stably transformed transgenic plant cell - useful for improving plant disease and pest resistance, and grain quality

This sequence is used to describe a method for regulating gene expression in a stably transformed transgenic plant cell. The method comprises controlled to chimeric genes (5' to 3') into the plant cell genome. The first gene comprises a promoter operably linked to a Gal4 binding complementary sequences), which is itself linked to a polyadenylation signal sequence. The Gal4 sequence is located upstream of the promoter; if a minimal promoter is used. The second chimeric gene comprises a promoter; and a DNA sequence encoding a companiant of Gal4 transcriptional activator, which is operably linked to a DNA sequence encoding a transcriptional activator and signal sequence of the first operably linked to a polyadenylation signal companies. The sequence is itself operably linked to a polyadenylation signal expression of the first chimeric gene regulates the companies of plant disease and pest resistance, in addition to grain quality (e.g. plant disease and pest resistance, in addition to grain quality (e.g. transcription factor) achieves a higher level of expression when compared to the level using the highly expressed seed storage protein gene compared to the level using the highly expressed seed storage protein gene compared complementally maintaining the expression pattern of a tissue-specific or Gaps ö Query Match

2.8%; Score 11.8; DB 1; Length 17;
Best Local Similarity 86.7%; Pred. No. 5e+02;
Matches 13; Conservative 0; Mismatches 2; Indels Sequence 17 BP; 1 A; 9 C; 5 G; 2 T; 0 U; 0 Other; developmentally regulated promoter 272 GGAGCAGGCGGCAC 286 16 gaagcagrecededec 2 ò

RESULT 1008

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AAA33417 standard; DNA; 17 BP. AAA33417

AAA33417;

28-JUL-2000 (first entry)

Low adenosine antisense oligonucleotide SEQ ID NO:1106.

Human; adenosine receptor; low adenosine antisense oligonucleotide; phosphorothioate; impaired respiration; inflammation; allergy; allergy; allergy; disease; bronchoconstriction; inhibitor; antinflammatory; antiallergic; antiaethmatic; cytostatic; analgesic; impaired alrway; lung disease; ischaemic condition; pulmonary visoconstriction; asthma; respiratory distress syndrome; pain; cystic fibrosis; emphysema; pulmonary thypertension; chronic obstructive pulmonary disease; COPD; cancer; leukaemia; lymphoma; carcinoma; metastasis; ss.

Homo sapiens

40200009525-A2.

24-FEB-2000.

99WO-US017712. 03-AUG-1999; 98US-0095212P. 03-AUG-1998;

(UYEC-) UNIV EAST CAROLINA.

Ayce JW;

WPI; 2000-205971/18.

New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstruction, inflammation, allergies, asthma, hypertension,

bronchitis, emphysema, respiratory distress syndrome, ischemia or cancers.

Claim 18; Page 403; 1343pp; English.

The present invention describes a new composition comprising an antisense oligonucleotide (ON) with low adenosine (up to 15%), which targets uncleic acids involved in bronchoconstriction, allergies, and/or inflammation. The ON can have antiniflammatory, antiallergic, antialsthmatic, cytostatic and analgesic activities. The compositions are impaired airways, including lung disease associated with inflammation, impaired airways, including lung disease associated with inflammation, impaired airways, including lung disease and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating conficiency pulmonary wasoconstriction, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, emphysema, chronic obstructive fibrosis, pulmonary disease (CODP), and cancers such as leukamias, lymphomas, carcinomas, and cancers which may metastasise to the lungs, including breast and prostate cancer. The reduction of the adenosine content of the cases of deoxyadenosine which may metastasise to the lungs, including controlle sequences side effects. The A-containing ONS break down with the crease of deoxyadenosine which activates adenosine receptors causing conchoconstriction and inflammation. AAA32313 to AAA3532 represent to invention, which correspond to SEQ ID NO:1 to 2815, and then the last 185 sequences are also called SEQ ID NO:1 to 185, but the sequences differ from the previously named sequences. SEQ ID NO:1 to 185, but the sequence differ contemporation of the present invention. N action with their corresponding SEQ ID NO: sequences given in the disclosure of the present invention do not match lighting.

Sequence 17 BP; 0 A; 7 C; 3 G; 7 T; 0 U; 0 Other;

0; Gaps Similarity 86.7%; Pred. No. 5e+02; 13; Conservative 0; Mismatches 2; Indels Query Match

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RESULT 1009 AAF19539

AAF19539 standard; DNA; 17 BP AAF19539;

Human IL3 receptor polynucleotide fragment #1106.

14-MAR-2001 (first entry)

human; airway disorder; bronchoconstriction; lung inflammation; human; airway disorder; bronchoconstriction; lung inflammation; human; airway disorder; bronchoconstriction; lung inflammatory; surfactant depletion; respiratory; bronchodilator; antifflammatory; immunosuppressive; antisathmatic; analgesic; hypotensive; cycostatic; respiratory obstruction; pulmonary obstruction; impeded respiration; surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory distress syndrome; pain; cystic fibrosis; allergic rhinitis; pulmonary hypertension; emphysems; pulmonary transplantation rejection; chronic obstructive pulmonary disease; pulmonary infection; bronchitis; cancer; ss.

Homo sapiens.

WO200062736-A2

26-OCT-2000

24-MAR-2000; 2000WO-US008020

99US-0127958P 06-APR-1999;

UNIV EAST CAROLINA. WPI; 2000-679539/66. (UYEC-) (Nyce JW;

Low adenosine (A) content antisense oligonucleotides which do not trigger adenosine receptors during metabolism, useful e.g. for treating cancers and respiratory obstructions.

Claim 14; Page 207; 1592pp; English.

The present invention describes low adenosine (A) content antisense oligomucleotides and compositions (I) comprising them. In the antisense oligomucleotides the A is replaced by a 'Universal' or alternative base. (C) can have respiratory, bronchodilator, antiinflammatory, analgesic, immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. The antisense oligomucleotides and (I) can be used to down-regulate the expression and or activity of target polypeptides associated with a ctivating peptide factors and transmitters, transcription factors, immunoslobulins and antibodies, antibody receptors, cytokines and chemokines, endogenously produced specific and non-specific enzymes, binding proteins, adhesion molecules and their receptors, cytokines and chemokine receptors, bradykinin receptors, cytokines and nervous system (CNS) and peripheral nervous and non-nervous system peptide receptors, bradykinin receptors, central nervous some non-nervous system peptide cransmitters, defensins, growth factors, vasoactive peptides and cransmitters, defensians, growth factors, vasoactive peptides and cransmitters, defensians, growth factors, vasoactive peptides and cransmitters, defensians and malignancy associated proteins. The antisense oligomucleotides may be used in this way to treat disorders including proteins and malignancy associated proteins. The antisense oligomucleotides may be used in this way to treat disorders and or bronchoconstriction, allergic with a disease or condition selected from pulmonary vasociated with a disease or surfactant hypoproduction which are associated with a disease or surfactant hypoproduction which are associated processes (CPDS), pain, cystic fibrosis (CP), allergic rhinitis (AR) pulmonary disease or conditions engagements and antisense oligomucleotides used in the exemplification of the resemblicions properties of properties are oligomucleotides used in the exemplification of the resemblicions which a described in the exemplification of the resemblicions. the present invention

Sequence 17 BP; 0 A; 7 C; 3 G; 7 T; 0 U; 0 Other;

Gaps Query Match
2.8%; Score 11.8; DB 1; Length 17;
Best Local Similarity 86.7%; Pred. No. 5e+02;
Matches 13; Conservative 0; Mismatches 2; Indels 0;

ઠે 셤 RESULT 1010 AAA25624,

AAA25624 standard; DNA; 17 BP. AAA25624; 19-JUL-2000 (first entry)

Oestrogen receptor hammerhead ribozyme target sequence SEQ ID NO:2122.

Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease; anticancer; breast cancer; endometrium cancer; ss.

Homo sapiens

V09954459-A2

Bellon L;

19-APR-1999; 20-APR-1998; 23-JUN-1998;

28-OCT-1999

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The present invention describes nucleic acids (A) that interact stably with a target sequence and contain at least one phosphoro(di)thioate with a target sequence and contain at least one phosphoro(di)thioate included acid (A) that modulates expression of the oestrogen receptor contains acid (A) that modulates expression of the oestrogen receptor.

Contains a transforming cells ex vivo and implanting treated cells, or correlate the high selectivity for targeted RNA, (A) can also be used to correlate inhibition of gene expression with alterations in phenotype, correlate inhibition of gene expression with alterations in phenotype, correlate inhibition of gene expression with alterations in phenotype, correlate inhibition of gene expression with alterations in phenotype, correlate inhibition of gene expression with alterations in (A) improves considerable for RNA, in the same way that restriction endonucleases are used with DNA). The combination of modifications in (A) improves to AAA2474 to nucleases, binding affinity and/or activity. AAA23503 to AAA2474 represent oestrogen receptor hardred ribozyme sequences.

Contains and AAA26105 represent their corresponding target sequences.

Contains and AAA26107 represent their corresponding target sequences.

Contains and AAA26107 represent their corresponding target and contains and antisense oligonucleotides used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                 New nucleic acids that interact, and optionally cleave, target sequences, used to treat cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oestrogen receptor hammerhead ribozyme target sequence SEQ ID NO:1302.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease; anticancer; breast cancer; endometrium cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.8%; Score 11.8; DB 1; Length 17;
86.7%; Pred. No. 5e+02;
artive 0; Mismatches 2; Indels
                                                                                                                                                                         Mcswiggen JA, Karpeisky A,
is T, Woolf T, Haeberli P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 3 A; 7 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                         Claim 77; Page 58; 148pp; English.
                                                                                                                                                                            Beigelman L, Mcswig
Zwick M, Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99WO-US008547.
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99WO-US008547.
                                              98US-0082404P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA24804 standard; DNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-JUL-2000 (first entry)
                                                                                                                          (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 86.74
These 13; Conservative
                                                                                                                                                                                                       Zwick M,
                                                                                                                                                                                                                                                                           WPI; 2000-013248/01.
                                                                                                                                                                                                                               Matulic-Adamic J;
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                                                                                                                                                                            Thompson JD,
Reynolds M,
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  19-APR-1999;
                                                   20-APR-1998;
23-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA24804;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nvention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ð
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes nucleic acids (A) that interact stably with a target sequence and contain at least one phosphoro(di)thioate with a target sequence and contain at least one phosphoro(di)thioate utility, having endomuclease activity. (A), and more generally any catalytic nucleic acid (A) that modulates expression of the oestrogen receptor gene, are used to treat cancer (particularly of breast or endometrium), in vivo or by transforming cells ax vivo and implanting treated cells, or for other conditions associated with levels of oestrogen receptor. Secause of the high selectivity for targeted RNA, (A) can also be used to correlate inhibition of gene expression with alterations in phenotype, correlate inhibition of gene expression with alterations in phenotype, correlate (for RNA, in the same way that restriction endomucleases are used with DNA). The combination of theoreticinon in (A) improves the cesistance to nucleases, binding affinity and/or activity. AAA23503 to AAA26105 represent their corresponding target sequences, and AAA26105 represent their corresponding target sequences. AAA25293 to AAA26105 represent their corresponding target sequences sequences. AAA26205 represent oestrogen receptor hairpin inboxyme sequences and antisence oligonucleotides used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ô
                                                                                                                                                                                                                                                                                                                                                                                        nucleic acids that interact, and optionally cleave, target sequences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oestrogen receptor hammerhead ribozyme target sequence SEQ ID NO:1301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease; anticancer; breast cancer; endometrium cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Gaps
                                                                                                                                                                                                                                     Bellon L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.8%; Score 11.8; DB 1; Length 17; 86.7%; Pred. No. 5e+02; ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                             Thompson JD, Beigelman L, Mcswiggen JA, Karpeisky A,
Reynolds M, Zwick M, Jarvis T, Woolf T, Haeberli P;
Matulic-Adamic J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seguence 17 BP; 1 A; 10 C; 1 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 77; Page 85; 148pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA24803 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           201 TCGGTGAAAGCAGAG 215
                                                                                                          98US-0082404P.
                                                            99WO-US008547
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TGGGGGAAAGCAGAG 2
                                                                                                                                                                                      RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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Best Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                         used to treat cancer.
                                                                                                                                                                                                                                                                                                                                          WPI; 2000-013248/01.
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nvention

WO9954459-A2

28-OCT-1999

Homo sapiens

19-JUL-2000

AAA24803;

RESULT 1011 AAA24803

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with a target sequence and contain at least one phosphoro(di)thioated link, having endonuclease activity. (A), and more generally any catalytic mucleic acid (A) that modulates expression of the oestrogen receptor gene, are used to treat cancer (particularly of breast or endometrium), in vivo or by transforming cells ex vivo and implanting treated cells, or floo other conditions associated with levels of oestrogen receptor. Because of the high selectivity for targeted RNA, (A) can also be used to correlate inhibition of gene expression with alterations in phenotype, particularly for identification of therapeutic targets, and as research reagents (for RNA, in the same way that restriction endonucleases are used with DNA). The combination of modifications in (A) improver.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   resistance to nucleases, building affinity and/or activity. AAA23503 to AAA24747 represent oestrogen receptor hammerhead ribozyme sequences, and AAA24748 to AAA2592 represent their corresponding target sequences. AAA24789 to AAA26105 represent their corresponding target sequences. sequences, and AAA26107 to AAA26218 represent their corresponding target sequences. and AAA26219 to AAA26218 represent their corresponding target sequences oligonucleotides used in the exemplification of the present
                                                                                                                                                                                                                                            nucleic acids that interact, and optionally cleave, target sequences,
                                                                                                                                                                                                                                                                                                                                                         present invention describes nucleic acids (A) that interact stably
                                                                                                           Beigelman L, Mcswiggen JA, J
Zwick M, Jarvis T, Woolf T,
                                                                                                                                                                                                                                                                                                                  Claim 77; Page 58; 148pp; English.
98US-0082404P.
98US-00103636.
                                                                 (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                         used to treat cancer
                                                                                                                                                                                                     WPI; 2000-013248/01.
                                                                                                                                                           Matulic-Adamic J;
20-APR-1998;
23-JUN-1998;
                                                                                                           Thompson JD,
Reynolds M,
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Seguence 17 BP; 4 A; 6 C; 4 G; 3 T; 0 U; 0 Other;

nvention

Gaps ö Query Match

2.8%; Score 11.8; DB 1; Length 17;

Best Local Similarity 86.7%; Pred. No. 5e+02;

Matches 13; Conservative 0; Mismatches 2; Indels

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AAA25625 standard; DNA; 17 BP 19-JUL-2000 (first entry) AAA25625; RESULT 1013 AAA25625

Oestrogen receptor hammerhead ribozyme target sequence SEQ ID NO:2123

Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease; anticancer; breast cancer; endometrium cancer; ss.

Homo sapiens

WO9954459-A2

18-OCT-1999

99WO-US008547. 19-APR-1999;

98US-0082404P. 98US-00103636. 20-APR-1998; 23-JUN-1998;

(RIBO-) RIBOZYME PHARM INC.

Bellon L; Karpeisky A, Haeberli P; Beigelman L, Mcswiggen JA, Zwick M, Jarvis T, Woolf T, Thompson JD, Beig Reynolds M, Zwick Matulic-Adamic J;

WPI; 2000-013248/01.

Bellon L;

Karpeisky A, Haeberli P;

nucleic acids that interact, and optionally cleave, target sequences used to treat cancer.

Claim 77; Page 85; 148pp; English.

the target sequence and contain at least one phosphoro (di)thioate with a target sequence and contain at least one phosphoro (di)thioate link, having endonuclease activity. (A), and more generally any catalytic nucleic acid (A') that modulates expression of the oestrogen receptor, gene, are used to treat cancer (particularly of breast or endometrium), or or by transforming cells ex vivo and implanting treated cells, or for other conditions associated with levels of oestrogen receptor.

C for other conditions associated with levels of oestrogen receptor.

Because of the high selectivity for targeted RNA, (A) can also be used to correlate inhibition of gene expression with alterations in phenotype, correlate inhibition of gene expression with alterations in phenotype, correlate inhibition of mene expression with alterations in phenotype, correlates in the same way that restriction endonucleases are reaged with DNA). The combination of modifications in (A) improves used with DNA). The combination of modifications in (A) improves the selection of the same way that restriction endonucleases are used with DNA). The combination of modifications in (A) improves the selection of the same receptor hammerhead ribozyme sequences, and AAA24148 to AAA26107 to AAA26218 represent their corresponding target sequences and sequences. AAA26219 tepresent other ribozyme sequences and sequences and sequences and sequences and the exemplification of the present The present invention describes nucleic acids (A) that interact stably with a target sequence and contain at least one phosphoro(di)thioate invention

Sequence 17 BP; 1 A; 9 C; 1 G; 6 T; 0 U; 0 Other;

Gaps Query Match
2.8%; Score 11.8; DB 1; Length 17;
Best Local Similarity 86.7%; Pred. No. 5e+02;
Matches 13; Conservative 0; Mismatches 2; Indels 0;

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15 TGGGGGAAAGCAGAG 1

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AAC70195 standard; DNA; 17 09-FEB-2001 (first entry) AAC70195;

RESULT 1014

ВР.

Single nucleotide polymorphism PCR primer #17.

Single nucleotide polymorphism; SNP; human; genetic disease; disease susceptibility; cardiovascular system; endocrine system; neurological system; forensic testing; paternity testing; PCR primer; WO200058519-A2. Homo sapiens 05-OCT-2000

99US-0127248P. 30-MAR-2000; 2000WO-US008440 31-MAR-1999; (WHED) WHITEHEAD INST BIOMEDICAL RES. (AFFY-) AFFYMETRIX INC.

201 TCGGTGAAAGCAGAG 215

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                                                                                                                                                                                                                                 The present invention is concerned with a number of human single mucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diseases sees of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, achizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Single nucleotide polymorphism; SNP; human; genetic disease; disease susceptibility; cardiovascular system; endocrine system; neurological system; forensic testing; paternity testing; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis.
                                                                                                 Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention is concerned with a number of human single
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      Lander ES;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.8%; Score 11.8; DB 1; Length 17;
86.7%; Pred. No. 5e+02;
tive 0; Mismatches 2; Indels
  Ireland JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 3 A; 4 C; 7 G; 3 T; 0 U; 0 Other;
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Cargill M, Daley GQ,
Patil N, Sklar P;
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Patil N, Sklar P;
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                                                                                                                                                                                                 Claim 8; Fig 5; 214pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC70192 standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3 GGACCGTGAGCTGGC 17
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                                                           WPI; 2000-611722/58.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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  Altshuler D,
Lipshutz RJ,
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Local St.
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                                                                                                                                                                                                                                                                                                                                                                                              diseases
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nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAMTU Displacement Protein (CDF). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
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86.7%; Pred. No. 5e+02;
cive 0; Mismatches 2; Indels
                                                                                                                                                                                                    2.8%; Score 11.8; DB 1; Length 17;
86.7%; Pred. No. 5e+02;
ive 0; Mismatches 2; Indels
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                                                                                                                                                                  Sequence 17 BP; 3 A; 4 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Zwick M, Pavco P, Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hammerhead ribozyme substrate #3199.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 54; Page 130; 164pp; English.
                                                                                                                                                                                                                                                                              314 GGACCGCGTGCTGGC 328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-AFR-2000; 2000WO-US009721
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                                                                                                                                                                                                                                                                                                                3 GGACCGTGAGCTGGC 17
                                                                                                                                                                                                                                                                                                                                                                                                          AAF06942 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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Best Local Similarity 86.7
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                interferon alpha; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200061729-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF06942;
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                                                                                                                                diseases
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Homo sapiens.

19-OCT-2000,

16-FEB-2001

AAF02141;

RESULT 1017

12-APR-1999;

Blatt L,

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The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 orphan receptor, ERA3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoletin, granulocyte colony stimulating factor protein and interferon alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
                                                                                                                                                                                                                                                  Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 2.8%; Score 11.8; DB 1; Length 17; Best Local Similarity 86.7%; Pred. No. 5e+02; Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 2 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                              Blatt L, Zwick M, Pavco P, Mcswiggen J;
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                                                                                                                                                                                                                                                                                                                                         Claim 37; Page 65; 164pp; English.
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                                                      11-APR-2000; 2000WO-US009721.
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                                                                                               99US-0129390P
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                                                                                                                                       (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                   WPI; 2000-647423/62.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200061729-A2
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                                                                                               12-APR-1999;
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                 19-0CT-2000.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1019
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, BAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                         Ribozyme; erythropoietin; granulocyte colony stimulating factor;
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86.7%; Pred. No. 5e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 2 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hammerhead ribozyme substrate #437.
                                                                                                                                                                                               Hammerhead ribozyme substrate #436.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 37; Page 65; 164pp; English
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                                                                          AAF02141 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zwick M, Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                          11-APR-2000; 2000WO-US009721.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                               interferon alpha; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-647423/62.
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Homo sapiens.

AAF02142;

BXSXXXXXXXXXXXXXXX

RESULT 1018

AAF02142,

Query Match

Matches

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Gaps

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The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATH Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and interferon alpha

Sequence 17 BP; 1 A; 8 C; 4 G; 4 T; 0 U; 0 Other;

2.8%; Score 11.8; DB 1; Length 17; 86.7%; Pred. No. 5e+02; tive 0; Mismatches 2; Indels 0; Gaps Query Match Best Local Similarity 86.7% Matches 13, Conservative

348 CTGCTCTACAGCGAC 362 2 crecrerchecec 16 ò

ABK00045 standard; RNA; 17 BP

ABK00045;

12-MAR-2002 (first entry)

Human NOGO Hammerhead Ribozyme #45.

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; noctropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOG0; hammerhead ribozyme; DNAzyme; inDzyme; G-cleaver; amberzyme; inzyme; lowphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphomy; leukaemia; human immunodeficiency virus; HIY associated NHL; mantle-cell lymphoma; MC1; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.

Homo sapiens. Synthetic.

WO200159103-A2.

39-FEB-2001; 2001WO-US004273.

11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P.

RIBOZYME PHARM INC.

(BLAT/) BLATT L. (MCSW/) MCSWIGGEN J. (BLAT/)

(CHOW/) CHOWRIRA B M.

Blatt L, Mcswiggen J, Chowrira BM;

WPI; 2001-607195/69.

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Claim 88; Page 66; 200pp; English

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NOGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a nucleic acids may be enzymatic nucleic acid cleaving a an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a NYM motif) proposessing an NCH motif), a G-cleaver (cleaving RNA with a NYM wothf) proposessing an NCH motif). The CD20-targetting nucleic acid is used to cleaver RNA of CD20 in the presence of a divalent cation that is preferably Mg^2+. Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more theraptes. In particular, the CD20 targetting nucleic acid may be used to the call and treat a patient having a condition associated with implementation of CD20. The treatment may further comprise the use of one or more indepth of CD20. The treatment may further comprise the use of one or more leave and the condition associated NEL, mantle-cell lymphona (NEL), immunocytoma (IMC), small B-cell lymphona; ICC leukaemia, HIV (human immunodeficiency virus) associated NEL, mantle-cell lymphona (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphona; immune thrombocytopaenia, and inflammatory arthropathy. The NOGO ctargetting nucleic acid may be contacted with a cell to reduce NOGO. The treatment may further comprise the use of one or more conclet a divalent cation that is preferably Mg^2 + Furthermore, the nucleic acid may be contacted with a cell to reduce NOGO activity of the nucleic acid may be contacted with a cell to reduce NOGO activity of the contact central nervous system (CNS) injury and cerebrovascular accident (CNA, stroke). Alzheimer's disease, disease, disease, disease, ataxia, Huntington's disease, or entral entral electronic sequence is a hammerhead ribozyme of t

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Sequence 17 BP; 0 A; 12 C; 2 G; 0 T; 3 U; 0 Other;

0; Gaps 2.8%; Score 11.8; DB 1; Length 17; 86.7%; Pred. No. 5e+02; ive 0; Mismatches 2; Indels 2.8% Query Match Best Local Similarity 86.7% Matches 13, Conservative

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RESULT 1021 ABK00895/c

ABK00895 standard; RNA; 17 BP.

ABK00895;

12-MAR-2002 (first entry)

Human NOGO Inozyme #165.

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; noctropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAZyme; inozyme; declarer; aumerzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; cerebrovascular accident; cVA; Alzheimer's disease; multiple sclerosis; Parkinson's disease; ataxia; Huntingcon's disease; ataxia; Huntingcon's disease; cerebrovascular dystrophy; neurodegenerative disease.

Homo sapiens.

40200159103-A2

16-AUG-2001,

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09-FEB-2001; 2001WO-US004273
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11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P. RIBOZYME PHARM INC.

RIBO-)

(BLAT/) BLATT L. (MCSW/) MCSWIGGEN J. (CHOW/) CHOWRIRA B M.

Chowrira BM; Mcswiggen J, Blatt L,

WPI; 2001-607195/69.

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Claim 88; Page 80; 200pp; English.

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NOGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a nucleic acids an NCH motif) and NCH motif), a G-cleaver (cleaving RNA with an NTM motif) propersion and NCH motif), a G-cleaver (cleaving RNA with an NTM motif) propersion and moberzyme (cleaving RNA with an NGN with an NGN motif) propersion of CD20 in the presence of a divalent cation that is preferably MG²+. Furthermore, it may be contacted with a call to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or folicular NHL, lymphocytic leukaemia, HIV (human immunodeficiency virus) associated NHL, mantle-cell lymphoma (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, insed to cleave RNA of the NOGO gene in the presence of a divalent cation that is preferably Mg²+. Furthermore, the cargetting nucleic acid may be contacted with a cell to reduce NOGO gene in the call and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more call and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more call and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more call and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more call and treat a patient having a condition associated with the level of NOGO treat central nervous system (NOGO) injury and cerebrovascular accident treatment standard the invention of disease, muscu sequence is an inozyme of the invention

Sequence 17 BP; 0 A; 12 C; 2 G; 0 T; 3 U; 0 Other;

o; Gaps ; Match
Local Similarity 86.7%; Pred. No. Se+02;
es 13; Conservative 0; Mismatches 2; Indels Query Match Matches

143 GGCGGTGGAGGCCGG 157

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15 GGAGGGGGAGGCCGG 1

ABK01170 standard; RNA; 17 BP ABK01170; RESULT 1022 ABK01170, axxxax

12-MAR-2002 (first entry)

Human NOGO Inozyme #440.

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; anctropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOG0; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; inizyme; ilozyme; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; Muman immunodeficiency virus; HIV associated NHL; mantie-cell lymphoma; MCL; immunocytoma; INC; immune thrombocytopechia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.

Homo sapiens. Synthetic. WO200159103-A2.

16-AUG-2001.

09-FEB-2001; 2001WO-US004273.

11-FEB-2000; 2000US-0191797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P.

(RIBO-) RIBOZYME PHARM INC.

(BLAT/) BLATT L. (MCSW/) MCSWIGGEN J. (CHOW/) CHOWRIRA B M.

Blatt L, Mcswiggen J, Chowrira BM;

WPI; 2001-607195/69.

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Claim 88; Page 85; 200pp; English.

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a nucleic acids may be enzymatic nucleic acid cleaving a an RNA molecule DNAzyme) an Inozyme (an endolytic nucleic acid cleaving a NN with a NNW motif), a an amberzyme (cleaving RNA with an NGN with a NNW motif) proposessing an NCH motif), a cleaver (cleaving RNA with a NGN with a PNR motif).

The CD20 in the presence of advalent carion that is preferably Mg² +.

Purthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more therapies. It particular, the CD20 targetting nucleic acid may be used to for treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular nucleic leukaemia, HIV (human immunodeficiancy virus) associated NHL, armitle-cell lymphoma (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, corrected with a cell to reduce NOGO gene in the presence of a divalent cation that is preferably Mg²2+, Furthermore, the presence of a divalent cation that is preferably Mg²2+, Furthermore, the cell and treat a patient having a condition associated with the level of the cell and treat a patient having a condition associated with the level of the cell and treat a patient having a condition associated with the level of the cell and treat a patient having a condition associated with the level of the cell and treat a patient having a congesteting multiple sclerosis (MS), chemotherapy-induced neuropathy, amyortophic lateral sclerosis (MS), parkinson's disease, ataxia, Huringrow's disease, central restrophy, and/or other neurodegenerative of the restrict disease, ataxia, Huringrow's disease, central restrophy, and/or other n

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Gaps

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Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NGO0; hammerhead ribozyme; DNAzyme; Inozyme; G-leaver; amberzyme; zinzyme; I-ymphoma; non-Hodgkin's lymphoma; NHL; lymphoma; leukaemia; human immunodeficiency virus; HIV associated NHL; manthe-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; Parkinson's disease; ataxia; Huntingcon's disease; ceretzel at accident ansocial amyorical ansocial amyorical ansocial amyorical ansocial amyorical amyorical amyorical ansocial ansocial amyorical ansocial adversal ataxia; Huntingcon's disease; certzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.
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states which respond to the modulation of NOGO expression. The present
                                                                                ch 2.8%; Score 11.8; DB 1; Length 17; 1 Similarity 86.7%; Pred. No. 5e+02; 13; Conservative 0; Mismatches 2; Indels
                                                  Sequence 17 BP; 2 A; 7 C; 2 G; 0 T; 6 U; 0 Other;
                    sequence is an inozyme of the invention
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                                                                                                                                                                                                                                                                                    ABK00894 standard; RNA; 17 BP.
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28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
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                                                                                                                                                          286 CCAAGCTGGTGAAGG 300
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(BLAT/) BLATT L.
                                                                                                                                                                                          15 CAAAACTGGTGAAGG 1
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                                                                                                                                                                                                                                                                                                                                                                                        Human NOGO Inozyme #164.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Blatt L, Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BLATT L.
MCSWIGGEN J.
CHOWRIRA B M.
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Best Local S
                                                                                                                                                                                                                                                    RESULT 1023
                                                                                                          Best Loca
Matches
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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NOGO). The nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN motif) pr an amberzyme (cleaving RNA with an NYN motif) pr

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cc with a YGY motil). The CD20-targeting mucleic acid is used to Cleave KNA CC CC CC COLOR in the presence of a divalent cation that is preferably Mg^2+.

CC FURCHORING. It may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level cc for CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targeting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-thody. Immunodeficiancy virus) associated NHL, lymphocytic lymphoma (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, communia, man inflammatory arthropathy. The NOGO-to targetting nucleic acid is used to cleave RNA of the NOGO gene in the presence of a divalent cation that is preferably Mg^2+. Furthermore, the presence of a divalent cation that is preferably Mg^2+. Furthermore, the concleic acid may be contacted with a cell to reduce NOGO activity of the concleic acid may be contacted with a cell to reduce NOGO activity of the concleic acid may be contacted with a cell to reduce NOGO activity of the concleic may further comprise the use of one or more therapies. In particular, the NOGO-targetting nucleic acid may be used to therapies. In particular, the NOGO-targetting nucleic acid may be used to comparise the use of one or more treat central nervous system (CNS) injury and cerebrovascular accident (CNA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemotherapy-induced neuropathy, anyotrophic lateral sclerosis (MS), sequence is an inozyme of the invention of NOGO expression. The present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; gene therapy; adenogine deaminase deficiency; p51; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDRV2A; melanoma; APC; HBA1; HBA2; adenomatcus polyposis of the Colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismatch repair; MSH2; MSH6; hyperilpidaemia; apolipoprotein B; LDLR; familial hypercholesterolaemia; UGT1; syndrome; APP; PSBN1; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presenilin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic;
nucleic acid is used to cleave RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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2.8%; Score 11.8; DB 1; Length 17;
Best Local Similarity 86.7%; Pred. No. 5e+02;
Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 0 A; 12 C; 3 G; 0 T; 2 U; 0 Other;
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27-WAR-2000; 2000US-0192179P.
01-JUN-2000; 2000US-0208538P.
30-OCT-2000; 2000US-0244989P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (UYDE ) UNIV DELAWARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antilipemic; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200173002-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABA77649;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1024
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Claim 7; Page 73; 294pp; English.
                                                                                                                           ABA77650 standard; DNA; 17 BP.
                                                                                                     337 ACCAGGGCCGGCTGC 351
                                                                                                           2 ACCAGTGCAGGCTGC 16
                                                                                                                                        24-JAN-2002 (first entry)
   WPI; 2001-639230/73.
                                                                                                                                                                                  antilipemic; ss.
                                                                                                                                                                                        Homo sapiens.
                                                                                                                                  ABA77650;
                                                                                         Query Match
                                                                                                                     RESULT 1025
                                                                                               Matches
                                                                                                                         ABA77650/
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WO200173002-A2.

04-OCT-2001.

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The present invention provides single-stranded oligonucleotides which can be used for the taxgeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin.

The following genes: adenosine deaminase, p53, beta-globin.

The following nalpha locus of this, personder kinase inhibitor 2A (CDKN2A), APC, Pactor V, Pactor VIII, Factor IX, haemoglobin alpha locus (DKN2A), haemoglobin alpha locus 2 (HBAZ), MIH1, MSH2, MSH6,

THEMAL), haemoglobin alpha locus 2 (HBAZ), MIH1, MSH2, MSH6,

TOPESCHILL, THOSE can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemophila, hypercholesterolaemia, thalassaemia, sickle cell anaemia, hariburs and seasons. The present sequence is one of the colon and various syndromes. The present sequence is one of the gene correcting oligonucleotides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human, gene therapy, adenosine deaminase deficiency; p53; beta-globin; retinoblastoma, BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor ZA; CDRVZA, melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VI; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOS; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein B; LDLR; familial hypercholesterolaemia; UGT1; syndrome, APP; PSEN1; antisense; UDP-glucurcnosyltransferase; amyloid precursor protein; presenilin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic;
                                                                                                                     Oligonuclectide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Beta globin mutation correcting oligonucleotide SEQ ID NO: 491.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 2.8%; Score 11.8; DB 1; Length 17; Best Local Similarity 86.7%; Pred. No. 5e+02; Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 2 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                           Claim 7; Page 73; 294pp; English.
Gamper HB, Rice MC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABA77645 standard; DNA; 17 BP.
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27-MAR-2000; 2000US-019219P.
UTUN-2000; 2000US-02653BP.
30-OCT-2000; 2000US-0244989P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           337 ACCAGGGCCGGCTGC 351
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                                                              WPI; 2001-639230/73.
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                                                                                                                                                                             modification.
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   Kmiec EB,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human, gene therapy, adenosine deaminase deficiency; p53; beta-globin, retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDNVAA, melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VI; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein B; LDLR; familial hypercholesterolaemia; UGT1; syndrome; APP; PSEN1; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presenilin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic;
                                                                                 Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical modification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 3 A; 7 C; 5 G; 2 T; 0 U; 0 Other;
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27-MAR-2000; 2000US-0192179P.
01-JUN-2000; 2000US-0208538P.
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01-JUN-2000; 2000US-0208538P.
30-OCT-2000; 2000US-0244989P.
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0; Gaps

Kmiec EB, Gamper HB, Rice MC;

WPI; 2001-639230/73.

Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical modification.

Claim 7; Page 73; 294pp; English.

The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CTR, cyclin-dependent kinase inhibitor 2A (CDKN2A), APC, Pactor VII, Factor IX, haemoglobin alpha locus (UGTI), amyloid precursor voil, Pactor (LDLR), manaeqlobin alpha locus 2 (UGTI), amyloid precursor protein (APC), presenilin-1 (PSEN1) and precursor protein (APC), presenilin-1 (PSEN1) and cut as cancer, adenosine deaminase deficiency, cystic fibrosis, haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, hambariamer's alseases melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting oligonucleotides of the invention

Sequence 17 BP; 3 A; 6 C; 5 G; 3 T; 0 U; 0 Other;

0; Gaps Query Match 2.8%; Score 11.8; DB 1; Length 17; Best Local Similarity 86.7%; Pred. No. 5e+02; Matches 13; Conservative 0; Mismatches 2; Indels

RESULT 1027

7646/c ABA77646 standard; DNA; 17 BP.

ABA77646;

24-JAN-2002 (first entry)

Beta globin mutation correcting oligonucleotide SEQ ID NO: 492.

Human, gene therapy, adenosine deaminase deficiency, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CFTR, cystic fibrosis, cancer, Factor V, cyclin-dependent kinase inhibitor ZA, CDKNZA, melanoma, APC, HBA1, HBA2, adenomatous polyposis of the colon, Factor VI, Factor II, thrombosis, haemophilia, alpha thalassaemia, haemoglobin alpha locus I, MiH1, APOR, mismatch repair; MSH2, MSH6, hyperlipidaemia, apolipoprotein E, LDLR, familial hypercholesterolaemia, UGT1, syndrome, APP, PSRN1, antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presenilin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic, antilipemic; ss.

Homo sapiens.

WO200173002-A2.

04-OCT-2001

27-MAR-2001; 2001WO-US009761.

27-MAR-2000; 2000US-0192176P. 27-MAR-2000; 2000US-0192179P. 01-UTW-2000; 2000US-0208538P. 30-OCT-2000; 2000US-024989P.

(UYDE) UNIV DELAWARE

Rice MC; Gamper HB, Kmiec EB,

WPI; 2001-639230/73.

Oligonucleotide for targeted alterations of genetic sequences and for trating cystic fibrosis, comprises at least one mismatch and chemical modification.

Claim 7; Page 73; 294pp; English.

The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these Sequences are directed at the following genes: adenosition deaminase, p53, beta-aglobin, retinoblastoma, BRCAL, BRCAL, CTR, Cyclin-dependent kinase inhibitor 2A (CDRN2A), APC, Factor V, Factor VII, Factor IX, haemoglobin alpha locus 2 (HBA1), haemoglobin alpha locus 2 (HBA2), MA11, MB12, MB16, apolitoprotein E (APOE), LDL receptor (LDLR), uDP-glucuroneyltransferase (UGT1), amyloid precursor protein (APC), presentlin-1 (PSEN1) and cuch as cancer, adenosine deaminase deficiency, cystic fibrosis, haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, arious syndromes. The present sequence is one of the gene correcting various syndromes. The present sequence is one of the gene correcting oligonucleotides of the invention

Sequence 17 BP; 3 A; 5 C; 6 G; 3 T; 0 U; 0 Other;

0; Gaps 2.8%; Score 11.8; DB 1; Length 17; 86.7%; Pred. No. 5e+02; tive 0; Mismatches 2; Indels Se+02; Se+02; Thes 2; Indels Local Similarity 86.79 tes 13; Conservative Query Match Matches

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AAH24022 standard; DNA; 17 BP. RESULT 1028 AAH24022

29-AUG-2001 (first entry)

Yeast GAL1/GAL10 promoter UASgal site, SEQ ID NO:5.

UASgal site; cis-acting transcription control element; Gal4; Gal3; Gal80; stoichiometrically balanced expression; yeast; galactose-inducible expression; expression construct; promoter; GAL1; GAL10; ds.

Saccharomyces cerevisiae.

US6221630-B1.

24-APR-2001.

99US-00275680. 24-MAR-1999;

(PENN-) PENN STATE RES FOUND.

4PI; 2001-307557/32.

Expression construct for inducing and sustaining high level recombinant polypeptide production in yeast, comprises nucleic acids encoding a trans

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-acting transcription factor, selectable marker and yeast origin of
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Disclosure, Col 15, 22pp, English.

The invention relates to high copy number expression constructs for high comparises a nucleic acid sequence encoding a set of trans-acting ranscription factors, a nucleic acid encoding a set of trans-acting ranscription factors, a nucleic acid encoding a set of trans-acting ranscription factors, a nucleic acid encoding a set of encoding a veat selectable marker providing an inefficiently or efficiently selected phenotype, a nucleic acid encoding a yeast or bacterial origin of replication (orl), and a unique restriction site downstream of a promoter containing a cis-acting transcription control element that is regulated by the transcription factors which are encoded by the expression construct provides for galactose-inducible protein expression. Such constructs contain DNA encoding the transcription factors (als) Gals) and a UNSagl cis-acting control element within the promoter which drives expression of the invention factors (als) and (also), becomes a cating control element within the promoter which drives expression of are expressed in stoichiometrically-balanced amounts, which is, and constitutive transcription factor, and can become toxic to the cell. The constitutive transcription factor, and can become toxic to the cell. The constitutive transcription factor, and can become toxic to the cell. The constitutive pression of the gene of interest. The expression constructs provide of the gene of interest. The expression constructs frowed expression of a gene of interest (which can encode robust, high level expression of a gene of interest (which can encode construct of the catual UNSgal sites found within the promoters of various yeast galactose-inducible genes which may be used as the cis-acting control element in a galactose-inducible genes by expression construct of the invention and provided constructs. the invention

Sequence 17 BP; 1 A; 9 C; 5 G; 2 T; 0 U; 0 Other;

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Query Match 2.8%; Score 11.8; DB 1; Length 17;
Best Local Similarity 86.7%; Pred. No. 5e+02;
Matches 13; Conservative 0; Mismatches 2; Indels 0; Gaps
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272 GGAGCAGGGCGCAC 286 16 GGAGCAGTGCGGCGC 2 ઠે

ABNO6222 standard; DNA; 17 BP. 29-MAY-2002 (first entry) ABN06222; RESULT 1029

Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:6214.

Human, genome-derived myosin-like protein 1, GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens

WO200192524-A2.

06-DEC-2001.

25-MAY-2001; 2001WO-US016981 26-MAY-2000;

2000US-0234687P. 2000US-0236359P. 2000GB-00024263. 2001WO-US000661. 2001WO-US000662. 2001WO-US000663. 30-JAN-2001; 2 30-JAN-2001; 2 30-JAN-2001; 2 21-SEP-2000; 27-SEP-2000; 04-OCT-2000;

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 claim be used as probes to detect, characterise and quantify nucleic acids can be used as probes to detect, characterise and quantify nucleic acids in samples, as amplification substrates, to hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antiologies that specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specifically as the rapport or and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser describtion ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 proteins, and in vaccines or for replacement therapy. The proteins and in vaccines or for replacement therapy. The protein mascle disorders. hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and selected disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence and the exemplification of the present invention. N.B. The present sequence data for this patent did not form part of the printed sequence data for this patent did not form part of the printed sequence data for this patent did not form part of the printed sequence data for this patent did not form part of the printed sequence data for this patent did not form part of the printed sequence data for this patent did not form part of the printed sequence data for this patent did not form part of the printed sequence or the exemplication of the printed sequence or the patent did not form part of the printed sequence or the patent did not form part of the printed sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                      Chen W,
                                                                                                                                                                                                                                                                                                                                      Gu Y, Ji Y, Penn SG, Hanzel DK, Rank DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; SEQ ID NO 6214; 214pp; English.
                                                                                                                 30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000659.
30-JAN-2001; 2001WO-US000670.
3FEB-2001; 2001US-0266860P.
                                                                                      2001WO-US000667
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Gaps ö Query Match 2.8%; Score 11.8; DB 1; Length 17; Best Local Similarity 86.7%; Pred. No. 5e+02; Matches 13; Conservative 0; Mismatches 2; Indels 291 CTGGTGAAGGACCTG 305

Seguence 17 BP; 4 A; 6 C; 4 G; 3 T; 0 U; 0 Other;

16 crerrecaddacere 2 à

ABN07566 standard; DNA; 17 BP.

RESULT 1030 ABN07566 ABN07566;

Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:7558. 29-MAY-2002 (first entry)

Human, genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin, chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens.

WO200192524-A2

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used as probes to detect, characterise and quantity nucleic acids can be used as probes to detect, characterise and quantity nucleic acids can be used as probes to detect, characterise and quantity protein variants having desired phenotyplic improvements, and for provide initial substrates for the recombinant engineering of hGDMLP-1 captorin variants having desired phenotyplic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polymeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins as sendards in assays used to determine the concentration and invacines for hGDMLP proteins, as specifically as for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 proteins as according to hGDMLP-1 may be used for diagnosing a clasorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the horselent squence at the exemplification of the present invention. N.B.

The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO XXX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New polypeptide, for raising antibodies that recognize hGDMLF-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                Gu Y, Ji Y, Penn SG, Hanzel DK, Rank DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 7558; 214pp; English.
                                                                                                                                                     2001WO-US000661.
2001WO-US000662.
2001WO-US000663.
2001WO-US000664.
                                                                                            2000US-0234687P.
2000US-0236359P.
2000GB-00024263.
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                                                                                                                                                                                                                                                                                                                            30-JAN-2001; 2001WO-US000670.
05-FEB-2001; 2001US-0266860P.
                                      25-MAY-2001; 2001WO-US016981
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2001WO-US000666
                                                                                                                                                                                                                                                                            2001WO-US000667
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                                                                                                                                                                    30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
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30-JAN-2001;
30-JAN-2001;
                                                                              26-MAY-2000;
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                                                                                               21-SEP-2000;
                                                                                                               27-SEP-2000;
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Seguence 17 BP; 5 A; 3 C; 6 G; 3 T; 0 U; 0 Other;

Gaps .. 0 / Match 2.8%; Score 11.8; DB 1; Length 17; Local Similarity 86.7%; Pred. No. 5e+02; nes 13; Conservative 0; Mismatches 2; Indels Query Match

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385 ACGACGCCCCAAGA 399 3 Arcacceccaaca 17 ઠે g

ABNOS995/C-ID ABNOS995 standard, DNA, 17 BP. XX ABNOS995; XX 29-MAX-2002 (first entry) RESULT 1031

Human, genome-derived myosin-like protein 1, GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss. Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:5987.

Homo sapiens.

WO200192524-A2.

06-DEC-2001.

25-MAY-2001; 2001WO-US016981

26-MAY-2000; 27-SEP-2000;

2001WO-US000666. 2001WO-US000667. 2001WO-US000661 001WO-US000663 2001WO-US000664 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 04-OCT-2000;

Chen W, Shannon ME;

30-JAN-2001; 2001WO-US000668 30-JAN-2001; 2001WO-US000669 30-JAN-2001; 2001WO-US000670 05-FEB-2001; 2001US-0266860P (AEOM-) AEOMICA INC. Gu Y, Ji Y, Penn SG, Hanzel DK, Rank DR, Chen W, Shannon ME; WPI; 2002-179446/23. New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.

Disclosure; SEQ ID NO 5987; 214pp; English.

The present invention describes a human genome-derived myosin-like protein 1 (AGDMLP-1). The protein and polynucleotide sequences of AGDMLP-1 close to be used as probes to detect, characteries and quantify nucleic acids in semples, as amplification substrates, to provide initial substrates for the recombinant engineering of AGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The HGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP to proteins, as standards in assays used to determine the concentration and/or amount specifically of HGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser describing in onisation, as the respective supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production and in vaccines or for replacement therapy. The production and shelters muscle disorders having appearing to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patient of the present invention. N.B. The sequence data for this patient of the printed in electronic format directly from WIPO and the formal and the formal of the present invention. at ftp.wipo.int/pub/published_pct_sequence

Sequence 17 BP; 5 A; 2 C; 6 G; 4 T; 0 U; 0 Other;

Gabe ö Query Match 2.8%; Score 11.8; DB 1; Length 17; Best Local Similarity 86.7%; Pred. No. 5e+02; Matches 13; Conservative 0; Mismatches 2; Indels

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify harderic acids can samples, as amplification substrates to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specifically recognise capture probes for surface-enhanced laser description ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 protuction, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 2. The present sequence in the exemplification of the present invention. N.B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                     Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart;
muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease;
skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                     Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:10468.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hanzel DK, Rank DR, Chen W, Shannon ME;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; SEQ ID NO 10468; 214pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-0CT-2000; 2000GB-00024263; 30-JAN-2001; 2001W0-US000661. 30-JAN-2001; 2001W0-US000662. 30-JAN-2001; 2001W0-US000664. 30-JAN-2001; 2001W0-US000666. 30-JAN-2001; 2001W0-US000666. 30-JAN-2001; 2001W0-US000666. 30-JAN-2001; 2001W0-US000666. 30-JAN-2001; 2001W0-US000666. 30-JAN-2001; 2001W0-US0006669. 30-JAN-2001; 2001W0-US0006669.
                                                                                                                  ABN10476 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAY-2001; 2001WO-US016981.
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2000US-0236359P.
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17 ACATGGACTTCCTCA 3
                                                                                                                                                                                              (first entry)
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                                                                                                                                                      ABN10476;
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human, genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1, heart; muscle; myosin, chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                  Gaps
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                                                                                                            Query Match 2.8%; Score 11.8; DB 1; Length 17; Best Local Similarity 86.7%; Pred. No. 5e+02; Matches 13; Conservative 0; Mismatches 2; Indels
                                                                             Sequence 17 BP; 1 A; 3 C; 10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure, SEQ ID NO 7564; 214pp; English.
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27. SEP-2000; 2000GS-0234687P.
27. SEP-2000; 2000GS-0234687P.
30. JAN-2001; 2001WO-US000661.
30. JAN-2001; 2001WO-US000663.
30. JAN-2001; 2001WO-US000666.
30. JAN-2001; 2001WO-US000667.
30. JAN-2001; 2001WO-US000669.
                                                                                                                                                                                                                                                                                                                ABN07572 standard; DNA; 17 BP.
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                                                                                                                                                                                       16 TGCGGGTGACCGAGG 30
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used as immunogens to raise antibodies that specifically recognise hGDMLP proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 therapeutic supplement in patients having specific deficiency in hGDMLP-1 polymuclection, and in vaccines or for replacement therapy. The polymuclectics are an in vaccines or for replacement therapy. The polymuclectics and in vaccines encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1; is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/published_pct_sequence
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Sequence 17 BP; 5 A; 3 C; 8 G; 1 T; 0 U; 0 Other;

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Query Match
2.8%; Score 11.8; DB 1; Length 17;
Best Local Similarity 86.7%; Pred. No. 5e+02;
Matches 13; Conservative 0; Mismatches 2; Indels 0; Gaps
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ABN08151 standard; DNA; 17 BP. 29-MAY-2002 (first entry) ABN08151; RESULT 1034

Human, genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss. Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:8143.

Homo sapiens.

WO200192524-A2

06-DEC-2001.

25-MAY-2001; 2001WO-US016981

2000US-0234687P 2000US-0236359P 2000GB-00024263 26-MAY-2000; 21-SEP-2000; 27-SEP-2000; 04-OCT-2000;

2001WO-US000661

2001WO-US000663 2001WO-US000664 2001WO-US000665 2001WO-US000666 2001WO-US000668 2001WO-US000669 2001WO-US000670 2001US-0266860P 2001WO-US000662 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001;

(AEOM-) AEOMICA INC

Chen W, Shannon ME; Ji Y, Penn SG, Hanzel DK, Rank DR, WPI; 2002-179446/23. Gu Y,

New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser descrption ionization, comprises human myosin-like protein hGDMLP-1.

Disclosure; SEQ ID NO 8143; 214pp; English.

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used as probes to detect, characterise and quantify nucleic acids can be used as probes to detect, characterise and quantify candous. In the hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polyspetides may be used as immunogens to raise antiodies that specifically recognise hGDMLP-1 proteins or polyspetides may be used as immunogens to raise antiodies that specifically recognise hGDMLP-1 proteins, as specifically as the reapeutic supplement in patients having specific deficiency in hGDMLP-1 proteins, as specific deficiency in hGDMLP-1 proteins as a man and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The proteins associated with the expression of hGDMLP-1 in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the horsented at the the examplification of the present invention. N.B. The sequence data for this patent did not form part of the printed cat from with was obtained in electronic format directly from WIPO at they was obtained in electronic format directly from WIPO at they was obtained in electronic format directly from WIPO at they was obtained in electronic format directly from WIPO at they was obtained in electronic format directly from WIPO at they was obtained in electronic format directly from WIPO at they was obtained in electronic format directly from WIPO at they was obtained in electronic format directly from WIPO at they was obtained in electronic format directly from WIPO at the present and they was obtained in electronic format

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Sequence 17 BP; 6 A; 3 C; 6 G; 2 T; 0 U; 0 Other;

0; Gaps 2.8%; Score 11.8; DB 1; Length 17; 86.7%; Pred. No. 5e+02; cive 0; Mismatches 2; Indels Best Local Similarity 86.7 Matches 13, Conservative Query Match

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ABN10475 standard; DNA; 17 BP. ABN10475; RESULT 1035

Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:10467.

(first entry)

29-MAY-2002

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens.

WO200192524-A2

06-DEC-2001.

2000US-0234687P, 2000US-0236359P, 04-OCT-2000; 2000GB-00024263. 30-JAN-2001; 2001WO-US000661. 30-JAN-2001; 2001WO-US000662. 25-MAY-2001; 2001WO-US016981 26-MAY-2000; 2000US-0207456P 21-SEP-2000; 27-SEP-2000;

30-JAN-2001; 2001WO-US000663. 30-JAN-2001; 2001WO-US000664. 30-JAN-2001; 2001WO-US000665. 30-JAN-2001; 2001WO-US0006657. 30-JAN-2001; 2001WO-US000669. 2001WO-US000668 30-JAN-2001;

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The present invention describes a human genome-derived myosin-like protein 1 (hdDMLP-1). The protein and polymuclectide sequences of hdDMLP-1 can be used in gene therapy and vaccine production. The hdDMLP-1 can be used as probes to detect, characterise and quantify nucleic acids can be used as probes to detect, characterise and quantify hdDMLP-1 nucleic acids in samples, as amplification substrates, to protein variants having desired phenotypic improvements, and for protein variants having desired phenotypic improvements, and for expressing the proteins. The hdDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hdDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hdDMLP proteins, as specifically according the capture probes for surface-enhanced laser describing as the concentration and in vaccines or for replacement therapy. The polymucleotide sequences encoding hdDMLP-1 may be used for diagnosing a disorder associated with the expression of hdDMLP-1, in particular heart and in vaccines of for replacement sequence represents an oligomer used in the screening of the hdDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not format directly from WIPO at fire, wipo.int/pub/published_pot_sequence.
                                                                                                                                                                                                 New polypeptide, for raising antibodies that recognize hCDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                    Chen W, Shannon ME;
                                                                                                    Hanzel DK, Rank DR,
                                                                                                                                                                                                                                                                                                         Disclosure; SEQ ID NO 10467; 214pp; English.
05-FEB-2001; 2001US-0266860P.
                                                                                                    Gu Y, Ji Y, Penn SG,
                                                                                                                                                     WPI; 2002-179446/23.
                                                   (AEOM-) AEOMICA INC.
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0; Gaps Query Match 2.8%; Score 11.8; DB 1; Length 17; Best Local Similarity 86.7%; Pred. No. 5e+02; Matches 13; Conservative 0; Mismatches 2; Indels Sequence 17 BP; 1 A; 3 C; 9 G; 4 T; 0 U; 0 Other;

16 TGCGGGTGACCGAGG 30 2 receedraAceared 16 ð d

ABN06001 standard; DNA; 17 BP. 29-MAY-2002 (first entry) ABN06001; RESULT 1036 ABN06001

Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:5993.

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

WO200192524-A2. Homo sapiens.

06-DEC-2001.

26-MAY-2000; 2000US-0207456P. 21-SEP-2000; 2000US-0234687P. 27-SEP-2000; 2000US-0236359P. 25-MAY-2001; 2001WO-US016981.

Chen W, Gu Y, Ji Y, Penn SG, Hanzel DK, Rank DR, 30-JAN-2001; 2001W0-US000663. 30-JAN-2001; 2001W0-US000664. 30-JAN-2001; 2001W0-US000665. 30-JAN-2001; 2001W0-US000666. 30-JAN-2001; 2001W0-US000667. 30-JAN-2001; 2001W0-US000669. 30-JAN-2001; 2001W0-US000669. 30-JAN-2001; 2001W0-US000669. 30-JAN-2001; 2001WO-US000661. (AEOM-) AEOMICA INC.

New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1. Shannon ME; WPI; 2002-179446/23.

Disclosure, SEQ ID NO 5993; 214pp; English.

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 uncleic acids in samples, as amplification substrates and quantify hGDMLP-1 uncleic acids in samples, as amplification substrates and quantify hGDMLP-1 uncleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specific blomclecule and/or amount specifically for hGDMLP proteins, as specific blomclecule and/or amount specifically recognise hGDMLP-1 and/or amount specifically recognise hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production and skeletal muscle disorders. hGDMLP-1 may be used for diagnosing a disorder aspeciate with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed the fuguration, but was obtained in electronic format directly from WIPO at fig. halpo.int/pub/published_pot_sequence

Gaps ; 0 Query Match
2.8%; Score 11.8; DB 1; Length 17;
Best Local Similarity 86.7%; Pred. No. 5e+02;
Matches 13; Conservative 0; Mismatches 2; Indels Sequence 17 BP; 5 A; 2 C; 7 G; 3 T; 0 U; 0 Other;

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351 CTCTACAGCGACTIC 365 15 CICIACAIGGACTIC 1 ઠે 셤

ABN08152 standard; DNA; 17 BP. 29-MAY-2002 (first entry) ABN08152; ABN08152

RESULT 1037

Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:8144.

Human, genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

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The present invention describes a human genome-derived myosin-like protein 1 (InDAMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used an gene therapy and vaccine production. The hGDMLP-1 can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids an samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having dealred phenotyplo improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specifically secondaries and for and/or amount specifically of hGDMLP-1 proteins, as specific blomolecule and/or amount specifically of hGDMLP-1 proteins, as specific blomolecule capture probes for surface-enhanced laser describing in hGDMLP-1 proteins and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The polynuclectide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 2.

The present sequence represents an oligomer used in the screening of the GDMLP-1 sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at fite, wipo.int/pub/published_pot_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gu Y, Ji Y, Penn SG, Hanzel DK, Rank DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; SEQ ID NO 8144; 214pp; English
                                                                                                                                                                                 2000US-0234687P.
2000US-0236359P.
2000GB-00024263.
2001WO-US000661.
2001WO-US000662.
2001WO-US000663.
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30-JAN-2001; 2001WO-US000668
30-JAN-2001; 2001WO-US000669
30-JAN-2001; 2001WO-US000670
05-FEB-2001; 2001US-0266860P
                                                                                                                                                                                                                                                                                                       30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
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                                       WO200192524-A2.
  Homo sapiens.
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. Match 2.8%; Score 11.8; DB 1; Length 17; Local Similarity 86.7%; Pred. No. Se+02; les 13; Conservative 0; Mismatches 2; Indels
 Query Match
                     Best Loca
Matches
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265 TGCACCTGGAGCAGG 279
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RESULT 1038 ABN06469 ID ABN06469 standard; DNA; 17 BP.

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Human, genome-derived myosin-like protein 1, GDMLP-1, hGDMLP-1, heart, muscle, myosin, chromosome 22, gene therapy, vaccine, heart disease, skeletal muscle disorder, amplicon, screening, ss.
                                    Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:6461.
                                                                                                                                                            04-0CT-2000; 2000GB-0024263.
30-JNN-2001; 2001W0-US000661.
30-JNN-2001; 2001W0-US000663.
30-JNN-2001; 2001W0-US000663.
30-JNN-2001; 2001W0-US000665.
30-JNN-2001; 2001W0-US000665.
30-JNN-2001; 2001W0-US000666.
30-JNN-2001; 2001W0-US000669.
30-JNN-2001; 2001W0-US000669.
30-JNN-2001; 2001W0-US000669.
                                                                                                                         25-MAY-2001; 2001WO-US016981
                     (first entry)
                                                                                                                                                                                                                                                             (AEOM-) AEOMICA INC.
                                                                                             WO200192524-A2.
                                                                                Homo sapiens.
                                                                                                                                                21-SEP-2000;
27-SEP-2000;
                     29-MAY-2002
                                                                                                            06-DEC-2001.
       ABN06469;
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Chen W, Shannon ME; Hanzel DK, Rank DR, Gu Y, Ji Y, Penn SG, WPI; 2002-179446/23. New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.

Disclosure; SEQ ID NO 6461; 214pp; English.

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 can be used as probes to detect, characterise and quantify nucleic acids can be used as probes to detect, characterise and quantify compared in the used as probes to detect, characterise and quantify protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specifically as the rapecute grobes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The collowed associated with the expression of hGDMLP-1, in particular heart and issociated with the expression of hGDMLP-1, in particular heart and skeletal mascle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO.

The present become the control of the present invention. We see the figuration that was obtained in electronic format directly from WIPO can fire the presence of the presence of

Sequence 17 BP; 2 A; 5 C; 3 G; 7 T; 0 U; 0 Other;

Query Match

2.8%; Score 11.8; DB 1; Length 17;

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04-OCT-2000; 2000GB-00024263.
30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000662.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
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21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0236359P.
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        Pred. No. 5e+02;
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86.78; "1.
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30-JAN-2001, 2001WO-US000661.

30-JAN-2001, 2001WO-US000662.

30-JAN-2001, 2001WO-US000663.

30-JAN-2001, 2001WO-US000663.

30-JAN-2001, 2001WO-US000665.

30-JAN-2001, 2001WO-US000666.

30-JAN-2001, 2001WO-US000666.

30-JAN-2001, 2001WO-US000669.

30-JAN-2001, 2001WO-US000669.

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ABN06223 standard; DNA; 17 BP.
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21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0236359P.
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                        13; Conservative
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muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease;
skeletal muscle disorder; amplicon; screening; ss.
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CC homometer acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 provide initial substrates for the recombinant engineering of hGDMLP-1 expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration add/or amount specifically of hGDMLP-proteins, as specific blomolecule capture probes for surface-enhanced laser desorption ionisation, as the production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed the present invention. N.B. The sequence data for this patent did not form part of the printed the present invention. With the property of the printed in electronic format directly from WIPO
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Query Match 2.8%; Score 11.8; DB 1; Length 17; Best Local Similarity 86.7%; Pred. No. 5e+02; Matches 13; Conservative 0; Mismatches 2; Indels 191 TATCCACTGCTCGGT 205

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ABN01016 standard; DNA; 17 BP RESULT 1041 ABNO101

(first entry) 29-MAY-2002 ABN01016;

Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:1008.

Human, genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens.

WO200192524-A2.

06-DEC-2001.

25-MAY-2001; 2001WO-US016981

2000US-0234687P. 2000US-0236359P. 2000GB-00024263.

30-JAN-2001; 2001WO-US000661. 30-JAN-2001; 2001WO-US000662. 30-JAN-2001; 2001WO-US000663.

30-JAN-2001; 2001WO-US000665. 30-JAN-2001; 2001WO-US000664. 30-JAN-2001; 2001WO-US000665. 30-JAN-2001; 2001WO-US000666. 30-JAN-2001; 2001WO-US000669. 30-JAN-2001; 2001WO-US000670 05-FEB-2001; 2001US-0266860P 30-JAN-2001; 2001WO-US000667

(AEOM-) AEOMICA INC.

Shannon ME; 3 Chen Rank DR, Hanzel DK, Su Y, Ji Y, Penn SG, WPI; 2002-179446/23

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polymuclectide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 calcided acids in samples, as amplification substrates, to hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 proteins are having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be corporatine proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specific biomolecute and/or amount specificatly of hGDMLP-1 proteins, as specific biomolecute capture probes for surface-enhanced laser desorption ionisation, as to formate a seciated sequences encoding hGDMLP-1 may be used for diagnosing a polymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1 in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

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Sequence 17 BP; 8 A; 3 C; 6 G; 0 T; 0 U; 0 Other;

0; Gaps Query Match
2.8%; Score 11.8; DB 1; Length 17;
Best Local Similarity 86.7%; Pred. No. 5e+02;
Matches 13; Conservative 0; Mismatches 2; Indels

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ABN06470 standard; DNA; 17 BP. RESULT 1042 ABN06470

29-MAY-2002 (first entry)

ABN06470;

Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:6462.

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; 88.

Homo sapiens.

WO200192524-A2.

06-DEC-2001.

30-JAN-2001; 2001WO-US000661. 30-JAN-2001; 2001WO-US000662. 30-JAN-2001; 2001WO-US000663. 30-JAN-2001; 2001WO-US000664 30-JAN-2001; 2001WO-US000665 30-JAN-2001; 2001WO-US000666 25-MAY-2001; 2001WO-US016981 26-MAY-2000;

Shannon ME;

Chen W,

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86.7%; Pred. No. 5e+02;
tive 0; Mismatches 2; Indels
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30-JAN-2001; 2001WO-US000667.
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30-JAN-2001; 2001WO-US000669.
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids in samples, as amplification substrates of dantify hucheir acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specific biomolecule and/or amount specifically of hGDMLP-1 proteins, as specific biomolecule and/or amount specifically of hGDMLP-1 proteins, as specific biomolecule capture probes for surface-enhanced laser describing injoinsation, as production, and in vaccines or for replacement therapy. The production and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hospital production, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                           New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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iive 0; Mismatches 2; Indels
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30-JAN-2001, 2001WO-US000666.
30-JAN-2001, 2001WO-US000667.
30-JAN-2001, 2001WO-US000669.
30-JAN-2001, 2001WO-US000669.
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30-JAN-2001; 2001WO-US000662.
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Best Local Similarity 86.7
Matches 13; Conservative
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